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(54) Title: ASSAYS FOR LIGANDS FOR NUCLEAR RECEPTORS

(57) Abstract

The present invention includes nuclear receptor heterodimer and nuclear receptor-coactivator peptide assays for identifying ligands for nuclear receptors, utilizing scintillation proximity and fluorescence resonance energy transfer (FRET), and methods of using identified ligands.

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ASSAYS FOR LIGANDS FOR NUCLEAR RECEPTORS

Cross reference to Related Applications

- 5 This application claims benefit under 35 U.S.C.119(e) of provisional application 60/135,097.

Background of the Invention

Farnesoid X Receptor (FXR) is an orphan nuclear receptor initially identified from a rat liver cDNA library (8) that is most closely related to the insect ecdysone receptor. FXR is a member of the nuclear receptor family of ligand-activated transcription factors that includes receptors for the steroid, retinoid, and thyroid hormones (6). Northern and in situ analysis show that FXR is most abundantly expressed in the liver, intestine, kidney, and adrenal (8;22). FXR binds to DNA as a heterodimer with the 9-cis retinoic acid receptor RXR. The FXR/RXR heterodimer preferentially binds to response elements composed of two nuclear receptor half sites of the consensus AG(G/T)TCA organized as an inverted repeat and separated by a single nucleotide (IR-1 motif) (8). An early report showed that rat FXR is activated by micromolar concentrations of farnesoids such as farnesol and juvenile hormone (8). However, these compounds failed to activate the mouse and human FXR, leaving the nature of the endogenous FXR ligand in doubt. The present invention demonstrates that several naturally-occurring bile acids bind and activate FXR at physiological concentrations (see also recent publications 17; 19; 24). As discussed in the present invention, the bile acids that serve as FXR ligands include chenodeoxycholic acid (CDCA), deoxycholic acid (DCA), lithocholic acid (LCA), and the taurine and glycine conjugates of these bile acids.

Bile acids are cholesterol metabolites that are formed in the liver and secreted into the duodenum of the intestine, where they have important roles in

the solubilization and absorption of dietary lipids and vitamins. Most bile acids (~95%) are subsequently reabsorbed in the ileum and returned to the liver via the enterohepatic circulatory system. The conversion of cholesterol to bile acids in the liver is under feedback regulation: Bile acids down-regulate the transcription of cytochrome P450 7a (CYP7a), which encodes the enzyme that catalyzes the rate limiting step in bile acid biosynthesis. There are data to suggest that FXR is involved in the repression of CYP7a expression by bile acids, although the precise mechanism remains unclear (21). In the ileum, bile acids induce the expression of the intestinal bile acid binding protein (IBABP), a cytoplasmic protein which binds bile acids with high affinity and may be involved in their cellular uptake and trafficking. Two groups have now demonstrated that bile acids mediate their effects on IBABP expression through activation of FXR, which binds to an IR-1 type response element that is conserved in the human, rat, and mouse IBABP gene promoters (14; 17).

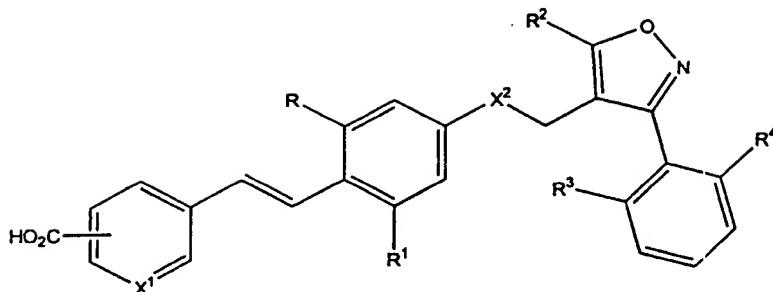
Thus FXR is involved in both the stimulation (IBABP) and the repression (CYP7a) of target genes involved in bile acid and cholesterol homeostasis.

The present invention has made the useful discovery of ligands for FXR as well as methods of determining genes regulated by FXR interaction and affecting those genes. Thus this discovery provides methods of regulating bile acid and cholesterol homeostasis, fatty acid absorption, and protein and carbohydrate digestion.

Summary of the Invention

The present invention provides a method of identifying compounds for the treatment of diseases or disorders modulated by FXR, comprising the step of determining whether the compound interacts directly with FXR, wherein a compound that interacts directly with FXR is a compound for the treatment.

The present invention also provides a compound that binds Farnesoid X receptor wherein the compound is of the following formula (II):



wherein X^1 is CH or N; X^2 is O or NH; R and R^1 may independently be H, lower alkyl, halogen, or CF_3 ; R^2 is lower alkyl; R^3 and R^4 may independently be H, lower alkyl, halogen, CF_3 , OH, O-alkyl, or O-polyhaloalkyl.

5 The present invention further provides a method of modulating a gene whose expression is regulated by FXR in a mammal comprising administering to the mammal a ligand of FXR.

Brief Description of the Drawings

10 **Fig. 1.** Shows CDCA selectively activates FXR as compared to other receptors. CV-1 cells were cotransfected with various nuclear receptor-GAL4 chimeras (25) and the reporter plasmid (UAS)₅-tk-CAT (26). Cells were treated with 100 μ M CDCA. Cell extracts were subsequently assayed for CAT activity (26). Data are expressed for each receptor as fold induction of CAT activity relative to vehicle-treated cells and represent the mean of three data points \pm SD.

20 **Fig. 2.** Shows activation of FXR by bile acids. (A) Chemical structures of major human bile acids. (B) Full-length human (filled bars) and full-length murine (open bars) FXR are activated by CDCA, LCA, and DCA. CAT assays were performed with extracts of CV-1 cells transfected with expression plasmids for human or murine FXR, human RXR α , and the FXREhsp27-tk-CAT reporter plasmid (27, 28). Cells were treated with 100 μ M of the indicated bile acid or farnesol, or 10 μ M of the indicated steroid or TTNPB. Cholesterol (CH).

25 (C to E) Dose response analysis of bile acids on human FXR. The assays

were run as above with 1, 3.3, 10, 33, or 100 μ M of CDCA, LCA, or DCA. (F) Human FXR is not activated by CDCA conjugates in CV-1 cells. The assay was run as above with 100 μ M free or conjugated CDCA. (G) Human FXR is activated by conjugated bile acids in CV-1 cells expressing the human IBAT gene. The assay was run as above except cells were additionally transfected with an expression plasmid for the human IBAT gene (pCMV-HISBT) and treated with 3 μ M of the indicated bile acid. Data are expressed as fold induction of CAT activity relative to vehicle-treated cells and represent the mean of three data points \pm SD (26).

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Fig. 3. Shows free and conjugated bile acids bind to FXR. (A) CDCA and its conjugates increase SRC1 binding to FXR. The FRET ligand-sensing assay was run with 10 nM of the biotinylated FXR LBD labeled with streptavidin-conjugated allophycocyanin, 10 nM of a SRC1 peptide labeled with streptavidin-conjugated europium chelate, and 100 μ M of the indicated compound. Data are expressed as the means \pm SD derived from three independent experiments. (B) Dose response analysis of CDCA binding to FXR. The FRET ligand-sensing assay was run in the presence of increasing concentrations of CDCA (open circles), glyco-CDCA (open triangle), tauro-CDCA (open boxes), or cholic acid (closed circles). (C) LCA, DGA, and TTNPB compete with CDCA for FXR binding. FRET ligand-sensing assays were run in the presence of 50 μ M CDCA and increasing concentrations of LCA (open circles), DCA (open boxes), and TTNPB (closed circles).

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Fig. 4. As shown in Fig. 4a, CDCA induced the interaction between FXR and the SRC-1 peptide. Dose response analysis showed that CDCA bound to FXR with an EC₅₀ value of ~4.5 μ M (Fig. 4b), which is well within the range of the intestinal concentration of this bile acid. Importantly, the closely related bile acids cholic acid and deoxycholic acid had no effect in the FXR FRET assay, demonstrating the specificity of CDCA binding to FXR (Fig. 4a).

30 Next examined was whether CDCA activated FXR in cells. CV-1 cells

were transiently transfected with an expression plasmid encoding the human FXR and a reporter construct containing the hsp27-EcRE, a response element that is known to be activated by the FXR/RXR heterodimer. A 130-fold activation of the FXR/RXR heterodimer could be achieved with 100 µM CDCA 5 (Fig. 4c).

Fig. 5 shows ligand binding to FXR measured by modulation of FXR:RXR heterodimer formation.

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Detailed Description of the Invention

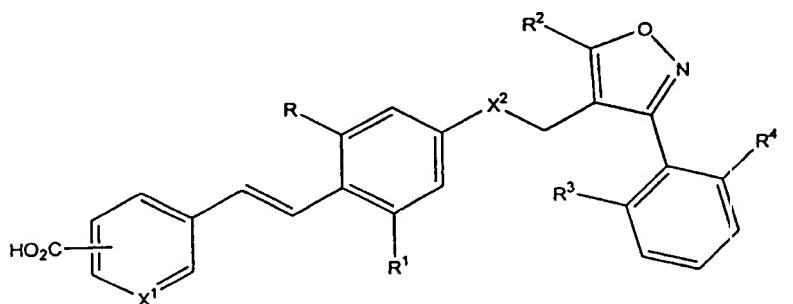
The present invention includes a generic approach to assay development for nuclear receptors, utilizing purified ligand binding domains. The concept of generic assay development has been extended to develop *in vitro* assays that 15 detect ligand binding by monitoring ligand induced changes in receptor heterodimerization. This approach has been demonstrated using both scintillation proximity and homogenous time-resolved fluorimetry (HTRF) in the accompanying examples but it is not restricted to these methods. Other marking and measuring techniques may also be used. However, the use of 20 scintillation proximity or HTRF provides a simpler and more practical methodology.

Another aspect of the invention is a nuclear receptor-peptide assay for identifying ligands. This assay utilizes fluorescence resonance energy transfer (FRET) and can be used to test whether putative ligands bind to FXR. The 25 FRET assay is based upon the principle that ligands induce conformational changes in nuclear receptors that facilitate interactions with coactivator proteins required for transcriptional activation.

In FRET, a fluorescent donor molecule transfers energy via a non-radiative dipole-dipole interaction to an acceptor molecule (which is usually a fluorescent 30 molecule). FRET is a standard spectroscopic technique for measuring distances in the 10-70 Å range. Upon energy transfer, which depends on the R^{-6} distance

between the donor and acceptor, the donor's fluorescence is reduced, and the acceptor fluorescence is increased, or sensitized. FRET is frequently used in both polymer science and structural biology and has recently been used to study macromolecular complexes of DNA, RNA, and proteins. In addition, Mathis has used europium cryptates with the multichromophoric Allophycocyanin to achieve an extremely large R_0 of 90 Å (29).

The present invention utilizes the above discussed assays for the discovery of ligands for FXR. Specifically, for example, the invention provides the discovery that CDCA binds and activates the farnesoid X receptor (FXR), an orphan member of the steroid hormone receptor superfamily which is activated by various farnesol metabolites and the synthetic retinoid TTNPB in cell-based assays. The invention additionally provides that a compound characterized by the following formula (I) is a ligand of FXR:

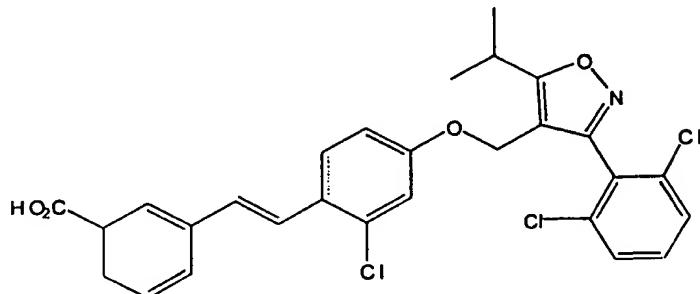


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wherein X^1 is CH or N; X^2 is O or NH; R and R^1 may independently be H, lower alkyl, halogen, or CF_3 ; R^2 is lower alkyl; R^3 and R^4 may independently be H, lower alkyl, halogen, CF_3 , OH, O-alkyl, or O-polyhaloalkyl.

An example of such a ligand is GW4064, a potent and selective FXR ligand, characterized by the following formula (II):

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Thus the present invention provides a method of regulating bile acid,
10 cholesterol, and lipid homeostasis using a compound, such as bile acid
(particularly CDCA) or a synthetic compound, such as GW4064, that binds the
FXR nuclear receptor.

Binding of the FXR nuclear receptor can result in the alteration of expression of various genes that FXR aids in regulating, including genes involved in lipid absorption and digestion in the small intestine and lipid homeostasis in the liver. Examples of such genes can include, but are not limited to, genes involved in bile acid transport, lipid absorption, cholesterol biosynthesis, proteolysis, amino acid metabolism, glucose biosynthesis, protein translation, electron transport, and hepatic fatty acid metabolism. It is noted that FXR often functions as a heterodimer with the Retinoid X Receptor (the FXR/RXR heterodimer). The inventive method includes using this technology to affect bile acid and cholesterol homeostasis such that, ultimately, cholesterol and lipid levels can be modified and in treating diseases in a mammal, including man, in which regulation of bile acid, cholesterol and lipid levels is important.

Thus the present invention provides a method of modulating a gene whose expression is regulated by FXR in a mammal comprising administering to the mammal a ligand of FXR: (a) Genes Downregulated in Liver, such as apolipoprotein B -2.7, plasma proteinase inhibitor alpha-1-inhibitor III group 3 (m22360) -2.7, L-gulono-gamma-lactone oxidase (d12754) -2.2, Peroxisomal enoyl-CoA:hydratase-3-hydroxyacyl-CoA bifunctional enzyme (k03249) -2.0,

liver fatty acid binding protein (L-FABP, m13501) -2.0, CYP4A2 (m57719) -1.9, CYP3A23 (x96721) -1.8
CYP3A1 (x64401) -1.6; (b) Genes Upregulated in Liver, such as small heterodimer partner homolog (d86580) +6.6, insulin-induced growth-response
5 protein, CL-6 (I13619) +5.9, elongation factor 2, EF-2 (y07504) +4.3, mouse cornichon +4.0, protein kinase C receptor (u03390) +3.1, mitochondrial cytochrome c oxidase (m27315) +2.7, cystathione gamma-lyase (x53460, d17370) +2.3, cytosolic phosphoenolpyruvate carboxykinase (k03243) +2.1, histidase (m58308) +2.1, S-adenosylmethionine synthetase (x60822) +2.0,
10 lanosterol 14-alpha-demethylase (u17697)+1.9, G protein-coupled purinoceptor P2U (I46865) +1.9, hepatic squalene synthetase (m95591) +1.5; (c) genes having altered expression in intestine: lipase (x61925) -9.5, pancreatic lipase (d88534) -4.4, colipase (m58370) -10.2, pancreatic phospholipase A-2 (d00036) -7.4, pancreatic amylase (m24962) -8.1, carboxypeptidase A1
15 (m23986) -6.1, carboxypeptidase A2 (m23721) -5.4, carboxypeptidase B (m23959) -4.2, pancreatic trypsin I (j00778) -6.8, pancreatic cationic trypsinogen (m16624), pancreatic trypsinogen II (v01274) -3.6, elastase I (v01234, I00112) -2.0, elastase II (I00118, I00124) -4.6, I-BABP (I22788) +2.3, intestinal fatty acid binding protein (FABP, k01180) -1.8, hepatic squalene
20 synthetase (m95591) +1.9, protein kinase C receptor (u03390) +1.6, elongation factor 2, EF-2 (y07504) +2.4, Small heterodimer partner homolog (d86580) +2.2. Additional genes can be determined by the methods taught herein.

By modulation can be meant upregulation or downregulation, alteration
25 in timing of expression or other means of modulation as known in the art. Modulation of these genes can be used to effect treatments of a variety of diseases and disorders, particularly those discussed herein.

In addition, it has been also found that CDCA, lithocholic acid (LCA), and
30 deoxycholic acid, (DCA) and their conjugated derivatives bind to FXR at concentrations consistent with those found in tissues and known to regulate gene transcription. These bile acids are highly efficacious activators of FXR in

cell-based reporter assays. With the conjugated forms of the bile acids, activation is only observed in cells that express a bile acid transporter. Thus, the conjugated derivatives, which account for ~98% of all bile acids in human bile, are likely to represent natural FXR ligands in tissues which express bile acid transporters whereas the unconjugated forms may function as ligands in tissues that do not express these transport proteins.

As can be seen from the examples below, including the studies regarding genes which expression is altered by FXR functioning, the present method can be used in a variety of treatments, including the use of FXR agonists to block fatty acid absorption in the intestine for the treatment of dyslipidemia (reduction of triglycerides, free fatty acids), obesity, and associated diseases (atherosclerosis); the use of FXR agonists to block protein and carbohydrate digestion in the intestine for the treatment of obesity; the use of FXR antagonists to block de novo cholesterol biosynthesis in the liver for the treatment of diseases related to elevated cholesterol levels, including atherosclerosis and gall stones; the use of FXR antagonists to block induction of SHP-1 expression (and consequent repression of CYP7A) for the treatment of diseases related to elevated cholesterol levels, including atherosclerosis and gall stones; the use of SHP-1 antagonists to block SHP-1-mediated repression of CYP7A for the treatment of diseases related to elevated cholesterol levels, including atherosclerosis and gall stones. Since FXR binds to DNA as an obligate heterodimer with RXR, the use of RXR-selective compounds (agonists or antagonists) can be made to effect any of the aforementioned effects via the FXR/RXR heterodimer.

In the assay of the present invention a method is provided for the rapid and simple determination of a ligand for a nuclear receptor which comprises contacting a component to be tested with an isolated nuclear receptor ligand binding domain which may be associated with a marking component, and a dimerization partner for the nuclear receptor ligand binding domain which is also associated with a marker; and measuring the interaction between the marking components to determine whether the component to be tested

modifies heterodimerization. Various known markers may be used in the process of the present invention such as radioactive markers. The marker could also be a fluorescent dye. When the marker is radioactive, scintillation proximity may be used to measure the marker. A suitable radioactive marker 5 would be radiolabelled GW4064 or derivative of formula (I). When the marker used is a fluorescent dye, homogenous time-resolved fluorimetry may be used to detect the marker. Other known marking and measuring techniques may be used depending on the marker. However, the markers need to be in close proximity to indicate heterodimerization. That is, to indicate that the component 10 to be tested functions as a ligand for the dimerization pair.

RXRs, such as RXR α , RXR β , and RXR γ , are examples of nuclear receptors that may be used as the dimerization partners in the process of the present invention. RXR α is exemplified in some of the examples. More dimerization partners may be known or later discovered which can readily be 15 utilized in the assay. It is preferable that the dimerization partners and the nuclear receptor ligand binding domains are recombinant proteins and preferably are bacterially expressed.

This method for the rapid determination of a ligand for a nuclear receptor comprises contacting a component to be tested with an isolated nuclear 20 receptor ligand binding domain which is associated with a first marking component, and a heterodimeric partner for the nuclear receptor ligand binding domain associated with a second marking component, and measuring the interaction between the marking components to determine whether the component to be tested modifies hetero-dimerization.

The first marking component may be a radioactive marker and the second marking component (or second marker) may be a SPA bead. The interaction of the markers in this case is determined by scintillation proximity. Alternatively, the first marking component may be a first fluorescent dye emitting at an emitting wavelength which excites the second marking 30 component which may be a second fluorescent dye. The interaction of the markers in this case is determined by homogenous time-resolved fluorimetry.

The interaction of the marking components in either case is measured by comparing a signal produced by a combination of the heterodimeric partner, the isolated nuclear receptor binding domain and the component to be tested with a signal produced by a combination of the heterodimeric partner and the
5 isolated nuclear receptor ligand binding domain in the absence of the component to be tested.

A compound of formula I or II can be detectably labelled by means known in the art. Radiolabelled GW4064 or a radiolabelled derivative of formula (I) can be used in a competition binding assay to detect compounds that directly
10 interact with FXR. Examples of competition binding assays include SPA assays where FXR is bound to an SPA bead in the presence of radiolabelled GW4064. Compounds that directly interact with FXR will displace radiolabelled GW4064 leading to a detectable signal (see Nichols, Anal Biochem. 1998).

15 Example 1

RXR α LBD

Human Retinoid X Receptor alpha Ligand Binding Domains:

RXR-alpha LBD (amino acids 225-462) was expressed in E. coli strain BL21 (DE3) as an amino-terminal polyHistidine tagged fusion protein. Expression
20 was under the control of an IPTG inducible T7 promoter. DNA encoding this recombinant protein and a modified histidine tag was subcloned into the expression vector pRSETa (Invitrogen). The sequence used in the construction of RXR-alpha LBD was derived from Genbank accession number X52773.

Ten-liter fermentation batches were grown in Rich P0₄ media with 0.1
25 mg/mL Ampicillin at 25°C for 12 hours, cooled to 9°C and held at that temperature for 36 hours to a density of OD₆₀₀ =14. At this cell density, 0.25 mM IPTG was added and induction proceeded for 24 hours at 9°C, to a final OD₆₀₀ =16. Cells were harvested by centrifugation (20 minutes, 3500g, 4°C), and concentrated cell slurries were stored in PBS at
30 -8°C.

Protein purification

Routinely, 40-50 g frozen cell paste (equivalent to 2-3 liters of the fermentation batch) was thawed and resuspended in 300mL TBS, pH 7.2, (25mM Tris, 150mM NaCl). Cells were lysed by three passages through a homogenizer (Rannie) and cell debris was removed by centrifugation (30 minutes, 20,000g, 5 °C). The cleared supernatant was filtered through coarse pre-filters and TBS, pH 7.2, containing 500 mM Imidazole was added to obtain a final imidazole concentration of 50 mM. This lysate was loaded onto a column (3 x 8 cm) packed with Sepharose [Ni⁺⁺ charged] Chelation resin (Pharmacia) and pre-equilibrated with TBS, pH 7.2, containing 50mM imidazole. After washing to baseline absorbance, the column was developed with a linear gradient of 50 to 500 mM imidazole in TBS, pH 7.2. Column fractions were pooled and dialyzed against TBS, pH 7.2, containing 5 mM DTT and .5mM EDTA. After dialysis the sample was concentrated using Centri-prep 10K (Amicon) and subjected to 10 size exclusion with a column (3 x 90 cm) packed with Sepharose S-75 resin (Pharmacia) pre-equilibrated with the same buffer.

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Biotinylation of Human Retinoid-X Receptor Ligand Binding Domain

Purified RXR α LBD biotinylation was carried out in a manner similar to 20 that described for FXR LBD in example 2.

Labeling of RXR α with CY5TM

Purified RXR α LBD was diluted to approximately 10 μ M in PBS and 25 approximately five-fold molar excess of Cy5TM monofunctional reactive dye [NHS ester] (Amersham Life Sciences) was added in a mininal volume of PBS. This solution was incubated in the dark with mixing for 30 minutes at ambient room temperature (approximately 23°C). The modification reaction was stopped by the addition of an excess of Tris-HCl, pH 8. Fluorescent dye modified RXR α LBD was dialyzed at 4°C, with minimal exposure to light, 30 against 4 buffer changes, each of at least 50 volumes, PBS containing 5mM DTT, 2mM EDTA, and 2% (w/v) sucrose. Aliquots were frozen on dry ice and

stored at -80°C.

Example 2

Determination of Ligand Binding to Farnasoid X Receptor: Retinoid X Receptor

5 Heterodimer utilizing Time Resolved Fluorimetry.

This example describes the use of ligand mediated heterodimerization to quantify ligand binding to the nuclear receptor Farnasoid X Receptor (FXR).

The method measures the ability of putative ligands to mediate the heterodimerization between the purified bacterial expressed FXR and RXR α

10 ligand binding domains (LBD). Detection of the associated LBD's are measured by time resolved fluorimetry (TRF). The purified LBD of FXR is labeled with biotin then mixed with stoichiometric amounts of europium labeled streptavidin (Wallac Inc). The purified LBD of RXR α is labeled with CY5TM.

Equimolar amounts of each modified LBD are mixed together and allowed to

15 equilibrate for at least 1 hour prior to the addition to either variable or constant concentrations of the sample for which the affinity is to be determined. After equilibration, the time-resolved fluorescent signal is quantitated using a fluorescent plate reader. The affinity of the test compound is estimated from a plot of fluorescence versus concentration of test compound added.

20 A basal level of FXR:RXR α heterodimer formation is observed in the absence of added ligand. Ligands that promote heterodimer formation induce a concentration-dependent increase in time-resolved fluorescent signal.

Compounds which bind equally well to both monomeric FXR and to the FXR:RXR α heterodimer would be expected to give no change in signal

25 whereas ligands which bind preferentially to the monomeric receptor would be expected to induce a concentration-dependent decrease in the observed signal.

METHODS & MATERIALS

Advance Preparation:

Human Farnasoid X Receptor alpha Ligand Binding Domain

5 Human FXR α Ligand Binding Domain (FXR α LBD) was expressed in E.coli strain BL21 (DE3) as an amino-terminal polyhistidine tagged fusion protein. Expression was under the control of an IPTG inducible T7 promoter. DNA encoding this recombinant protein was subcloned into the pRSET-A expression vector (Invitrogen). The coding sequence of Human FXR α LBD
10 was derived from Genbank accession number U 68233 (amino acids 222 to 472).

15 Ten-liter fermentation batches were grown in Rich P0₄ media with 0.1 mg/mL Ampicillin at 25°C for 12 hours, cooled to 9°C and held at that temperature for 36 hours to a density of OD₆₀₀ = 14. At this cell density, 0.25 mM IPTG was added and induction proceeded for 24 hours at 9°C, to a final OD₆₀₀ = 16. Cells were harvested by centrifugation (20 minutes, 3500g, 4°C), and concentrated cell slurries were stored in PBS at -8°C.

Purification of Receptor Ligand Binding Domain

20 Routinely, 30-40 g cell paste (equivalent to 2-3 liters of the fermentation batch) was resuspended in 200-250 mL TBS, pH 7.2 (25mM Tris, 150 mM NaCl). Cells were lysed by passing 3 times through a French Press and cell debris was removed by centrifugation (30 minutes, 20,000g, 4°C). The cleared supernatant was filtered through course pre-filters, and TBS, pH 7.2,
25 containing 500 mM imidazole was added to obtain a final imidazole concentration of 50mM. This lysate was loaded onto a column (6 x 8 cm) packed with Sepharose [Ni⁺⁺charged] Chelation resin (Pharmacia) and pre-equilibrated with TBS pH 7.2/ 50mM imidazole. After washing to baseline absorbance with equilibration buffer, the column was washed with one column
30 volume of TBS pH 7.2 containing 90mM imidazole. FXR α LBD was eluted directly with 365 mM imidazole. Column fractions were pooled and dialyzed

against TBS, pH 7.2, containing 0.5mM EDTA and 5mM DTT. The dialyzed protein sample was concentrated using Centri-prep 10 K (Amicon) and subjected to size exclusion, using a column (3 x 90 cm) packed with Sepharose S-75 resin (Pharmacia) pre-equilibrated with TBS, pH 7.2,
5 containing 0.5mM EDTA and 5mM DTT.

Biotinylation of FXR

Purified FXR α LBD was desalted/buffer exchanged using PD-10 gel filtration columns into PBS [100mM NaPhosphate, pH 7.2, 150mM NaCl].
10 FXR α LBD was diluted to approximately 10 μ M in PBS and five-fold molar excess of NHS-LC-Biotin (Pierce) was added in a minimal volume of PBS. This solution was incubated with gentle mixing for 30 minutes at room temperature. The biotinylation modification reaction was stopped by the addition of 2000x molar excess of Tris-HCl, pH 8. The modified FXR α LBD was dialyzed against
15 4 buffer changes, each of at least 50 volumes, PBS containing 5mM DTT, 2mM EDTA and 2% sucrose. The biotinylated FXR α LBD was subjected to mass spectrometric analysis to reveal the extent of modification by the biotinylation reagent. In general, approximately 95% of the protein had at least a single site of biotinylation; and the overall extent of biotinylation followed a
20 normal distribution of multiple sites, ranging from one to nine. —

RXR α LBD

RXR α LBD was prepared and labeled with CY5 TM in accordance with the procedures set forth in example 1.
25

Preparation of CY5 TM -RXR:Streptavidin-(Europium Chelate)-FXR Complex

Equimolar concentrations of biotinylated FXR and streptavidin-conjugated europium chelate were incubated in assay buffer containing 10 mM DTT for at least 10 minutes. To this solution an equimolar concentrations of
30 Cy5TM labeled RXR α was added and allowed to equilibrate for at least 30 min. The premixed receptor was then added in a one-step addition to the compound

plate, utilizing e.g., a Titertek Multidrop 384.

Materials:

Assay Buffer: 50 mM KCl, 0.1 mg/mL BSA, 10 mM DTT, and 50 mM Tris

(pH 8) The stock buffer is made by dissolving 2.853g Tris base, 4.167 g Tris

5 hydrochloride, 3.73 g KCl, and 0.1 g fatty acid free bovine serum albumin, in 1 L of deionized water. The pH is checked and adjusted to 8.0, if necessary, before adjusting to final volume. 0.154 g of solid DTT is added per 100 mL of buffer just before the start of an experiment.

BSA, fatty acid free

10 DTT

KCl

Europium labeled Streptavidin: (Wallac CR28-1 00)

Tris Hydrochloride

96 well plates: polypropylene for intermediate dilutions (Costar #3794)

15 and either a clear-bottomed white SPA plates (Costar #3632) or a black Polyfiltronics plate (UP350 PSB) for assays.

Methods:

Experimental Details:

20 Each well to be assayed contained a previously prepared solution of CY5TM RXR α and Europium labeled FXR and the desired concentration of test samples or controls (100 μ L total volume). In general, the total volume was held constant by varying the concentration and volume of premixed receptors to compensate for any changes in the volume of a particular set of samples. The
25 plates were incubated for at least 2 hours at room temperature and the fluorescent signal determined in a Wallac Victor Multilabel Fluorescence Reader.

Data Reduction:

For single concentration assays, the results of each test well were
30 expressed as % of control, C, calculated according to eq. 1.

$$C = 100 * \frac{F_{\text{sample}} - F_{\text{basal}}}{F_{\text{std}} - F_{\text{basal}}} \quad (1)$$

5 where F_{sample} is the signal observed in a particular sample well, F_{total} is the signal observed in the presence of control inhibitor and F_{basal} is the count rate observed in the presence of no ligand. The values used for F_{std} and F_{basal} were averages of the corresponding control wells included on every plate.

10 For Dose response assays, the data were first normalized to % of control using eq. (1). A plot of C_L , the % of control observed at ligand concentration L , versus ligand concentration, L was constructed. The data were fit to equation (2) to obtain best-fit parameters for the EC_{50} , F_{max} and F_{basal} .

$$15 C_L = F_{\text{basal}} + \frac{F_{\text{max}} * L}{EC_{50} + L} \quad (2)$$

Note that F_{max} , the maximal amplitude observed at saturating ligand concentrations, can be either a positive or negative value. The sign of this parameter indicates whether a particular test compound favors binding to the 20 FXR:RXR complex (positive F_{max}) or to either of the component receptors in a non-heterodimeric state (negative F_{max}). Furthermore, both F_{max} and F_{basal} are expressed in units of % of a standard compound.

RESULTS

25 Both the magnitude and sign of F_{max} (the maximal ligand-induced amplitude) must be considered. Note that the definition of the maximal response observed for CDCA = 100% is an arbitrary assignment. The purpose of the normalization is only to allow comparison of values obtained for different compounds on different days and/or using different fluorescence plate readers.
30 The results are shown in Figure 5.

Example 3

FXR is abundantly expressed in liver, intestine, kidney, and adrenal,

tissues through which there is significant bile acid flux. Based upon these observations and the FRET and transactivation data, the FXR represents a novel hormone signaling pathway and a transcription factor that induces the expression of I-BABP by CDCA. Thus, identification of ligands that bind to 5 FXR provides a method of effecting some of the therapeutic effects of CDCA through the regulation of bile acid flux, including desaturation of bile, inhibition of intestinal cholesterol absorption, and reduction in triglycerides.

In principle, the biology behind this method is to cause more bile acid to be transported through the gut of a mammal, including man. Thus, the FXR 10 receptor is a target for compounds (such as bile acid or other discovered ligands) that bind it and regulate bile acid transport.

Methods related to inhibiting binding of ligands, such as bile acid and GW4064, to FXR and to activating/inhibiting expression of intestinal bile acid-binding protein (I-BABP, which protein is activated by binding to bile acid-FXR- 15 RXR and is involved in transport of bile acid across membranes in the intestine) are included. Furthermore, increasing the transport of bile acids; stimulating the body to convert cholesterol into bile acid; reducing levels of cholesterol; inhibiting the intestinal absorption of cholesterol; reducing the levels of triglycerides; and treating diseases such as atherosclerosis, gallstone 20 disease, lipid storage disorders are also included. ——

FXR is a Nuclear Receptor for Chenodeoxycholic Acid

As noted supra, recently a cytosolic binding protein termed intestinal bile acid-binding protein (I-BABP) or fatty acid binding protein 6 (FABP6) was 25 identified as a component of the bile acid transport system in the ileal enterocyte. Kanda et al. showed that expression of I-BABP is dramatically induced by human bile, with chenodeoxycholic acid (CDCA) showing a significantly greater induction effect than other more hydrophilic bile acids. On the basis of this structural activity, Kanda proposed that an unidentified nuclear 30 receptor for CDCA was a transcriptional regulator of the expression of I-BABP in the intestine. It has now been discovered that CDCA binds and activates

FXR, an orphan member of the steroid hormone receptor superfamily.

To examine whether CDCA mediated its transcriptional effects through an orphan member of the steroid/retinoid/thyroid hormone receptor family (4) a chimeric receptor system in which the putative ligand binding domain (LBD) of 5 the human orphan receptor is fused to the DNA-binding domain of the yeast transcription factor GAL4 was used. Expression plasmids for the human nuclear receptor-GAL4 chimeras were generated by amplification of the cDNA encoding the putative LBDs and insertion into a modified pSG5 expression vector (Stratagene) containing the GAL4 DNA binding domain (amino acids 1 10 to 147) and the SV40 Tag nuclear localization signal (5). In CV-1 cells, CDCA selectively activated FXR [NR1H4] (Fig. 1), an orphan nuclear receptor expressed predominantly in the liver, kidney, intestine, and adrenals (7, 8). This strong activation by CDCA was unanticipated because FXR responds to high concentrations of farnesoids (7) and retinoids (9).

15 To further investigate the structure-activity relationship of FXR activation, a number of naturally occurring cholesterol metabolites, including bile acids (Fig. 2A), oxysterols and steroids, were tested for their ability to activate full-length human or full-length murine FXR in CV-1 cells. CDCA activated both the human and mouse FXR (Fig. 2B). Dose response analysis showed that 20 although some activation was seen at 3.3 μ M, greater activation was observed at 100 μ M CDCA (Fig. 2C). FXR was also activated by the secondary bile acids lithocholic acid (LCA) and deoxycholic acid (DCA), although these compounds were less efficacious than CDCA (Fig. 2B to E). A similar activation pattern was observed in various cell lines, including Drosophila-derived S2 cells, indicating that CDCA, LCA, and DCA do not require any 25 specialized metabolic conversion to activate FXR (11). Urso-DCA, the 7 β -hydroxy epimer of CDCA, and cholic acid (CA), which differs from CDCA by only the addition of a hydroxyl group at the 12 α position, were inactive on FXR (Fig. 2B). In addition, no activation of FXR was seen with either α - or β -30 muricholic acid (MCA), the glycine or taurine conjugates of bile acids, oxysterols, farnesol or other products derived from the mevalonate pathway

(Fig. 2B and F). Thus, both the 5 β -cholanoic acid backbone and stereochemistry of the hydroxyl groups in CDCA are critical for optimal FXR activation.

Bile acids are usually found conjugated to glycine or taurine, a derivative
5 of cysteine. Cells require the presence of an active bile acid transporter for uptake of these conjugated derivatives (12). To test whether conjugated bile acids would also activate FXR, the human ileal bile acid transporter (IBAT) was coexpressed with FXR in CV-1 cells (13). FXR was strongly activated by 3 μ M of the taurine or glycine conjugates of CDCA, LCA, and DCA (Fig. 2G).
10 Weaker activation was seen with the conjugated forms of CA and tauro-MCA was inactive (Fig. 2G). These data indicate that FXR can be activated by conjugated bile acids in tissues that express bile acid transporters such as the terminal ileum, liver, and kidney. The relationship between the chemical structure of bile acids and their activation of FXR is in close agreement with the
15 reported effects of bile acids on induction of I-BABP expression in Caco-2 cells and inhibition of CYP7a expression in hepatocytes (3,12)

Example 4

A fluorescence resonance energy transfer (FRET) assay as described further infra, was used to test whether putative ligands bound to a nuclear receptor, FXR. The FRET assay is based upon the principle that ligands
20 induce conformational changes in nuclear receptors that facilitate interactions with nuclear receptor coactivator proteins required for transcriptional activation. The ligand binding domain of FXR labeled with fluorophore allophycocyanin (APC) was incubated with a peptide containing the nuclear receptor interaction
25 domain from the SRC-1 coactivator labeled with europium cryptate.

Both the transactivation and ligand-binding assays typically used to determine activity of putative nuclear receptor activators assess the effect of test ligands on isolated receptor. However, a large proportion of the known orphan nuclear receptors interact with cofactor or coactivator proteins as a complex. Therefore, the present invention suggests the possibility that ligand binding to nuclear receptors may be modulated by the receptor's complexation
30

with a cofactor peptide. The ability of ligand to induce changes in the degree of this complex was then used as a basis for an inventive assay for the discovery of nuclear receptor ligands. Certain sequences of the cofactor may only be required to interact with the nuclear receptor. Various sequences of
5 the two cofactor proteins SRC-1 and CBP were synthesized and tested in HTRF and Biacore to determine the best sequences to use. The peptide, CPSSHSSLTERHKILHRLLQEGSPS-CONH₂ (SEQ ID NO.:1), i.e., SRC-1(LCD2,676-700) was used in screening efforts with FXR and this forms a further aspect of this invention.

10 Coactivator proteins interact with nuclear receptors in a ligand-dependent manner and augment transcription (9). A short amphipathic α -helical domain that includes the amino acid motif LXXLL (SEQ ID NO.:6) (L is Leu and X is any other amino acid) serves as the interaction interface between these coactivator molecules and the ligand-dependent activation function (AF-2)
15 located in the COOH-terminus of the nuclear receptor LBD (10). To test whether ligands would induce a conformation of FXR that favors coactivator binding, a cell-free ligand-sensing assay utilizing fluorescence resonance energy transfer (FRET) to monitor allosteric interaction of a peptide based on the sequence of the steroid receptor coactivator 1 (SRC1) with the receptor
20 was established. The use of FRET to monitor macromolecular complex formation is well established, particularly for immunoassays (11), and this detection methodology has recently been extended to characterize ligand binding to nuclear receptors (12).

Human FXR LBD was prepared and fluorescently labeled as described in
25 Example 2. The LBD of human FXR was labeled with the fluorophore allophycocyanin and incubated with a peptide derived from the second LXXLL (SEQ ID NO.: 6) motif of SRC1 (amino acids 676 to 700) that was labeled with europium chelate. The FRET ligand-sensing assay was performed by incubating 10 nM of the biotinylated FXR LBD that was labeled with
30 streptavidin-conjugated allophycocyanin (Molecular Probes) and 10 nM of the SRC1 peptide [amino acids 676 to 700, 5'-biotin-

CPSSHSSLTERHKILHRLQEGSPS-CONH₂] (SynPEP) (SEQ ID NO.:1) that was labeled with streptavidin-conjugated europium chelate (Wallac), in 50 mM Tris pH 8, 50 mM KCl, 0.1 mg/ml BSA, 1 mM EDTA, and 10 mM DTT, in the presence of test compound for 2 hours at 22°C. Data were collected using a

5 Wallac Victor™ fluorescence reader in a time-resolved mode. The relative fluorescence was measured at 665nM and the data reduction was as described in Example 1.

Preparation of Streptavidin-(Europium Chelate)-SRC1:Streptavidin-(APC)-FXR Complex

10 Biotinylated SRC-1(LDC2,676-700) peptide and a ½ stoichiometric amount of streptavidin-conjugated europium chelate were incubated in assay buffer containing 10mM DTT for at least 30 minutes. A second solution of stoichiometric amounts of biotinylated FXR and streptavidin-conjugated APC were incubated in assay buffer containing 10mM DTT for at least 30 minutes.

15 Each solution was then blocked with a 5 fold molar excess of biotin and allowed to equilibrate for at least 15 minutes. The labeled receptor and labeled peptide were mixed and again allowed to equilibrate for at least 30 minutes, then added in a one-step addition to the compound plate, utilizing, e.g., a Titertek Multidrop 384.

20 Materials:

APC-labeled streptavidin FXR and Europium labeled streptavidin SRC-1(LDC2,676-700)

SRC-1(LDC2,676-700):(SynPEP)

APC-Labeled streptavidin FXR and Europium labeled streptavidin

25 SRC1(LCD2,676-700)

Biotinylated Human Farnasoid-X receptor LBD:

Biotinylated SRC-1(LDC2,676-700):Biotin-CPSSHSSLTERHKILHRLL-QEGSPS-CONH₂ (SynPEP)

(SEQ ID NO.:1)

30 Assay Buffer: 50 mM KCl, 2mM EDTA, 0.1 mg/mL BSA, 10 mM DTT, and 50 mM Tris (pH 8). The stock buffer is made by dissolving 2.853g

Tris base, 4.167 g Tris hydrochloride, 3.73 g KCl, 0.74 g EDTA (disodium salt, dihydrate) and 0.1 g fatty acid free bovine serum albumin, in 1 L of deionized water. The pH is checked and adjusted to 8.0, if necessary, before adjusting to final volume. 0.154 g of solid DTT is added per 100 mL of buffer just before the start of an experiment.

5 BSA, fatty acid free

DTT

EDTA, disodium salt dihydrate

KCl

10 Allophycocyanin labeled streptavidin: (Molecular Probes S-868)

Europium labeled Streptavidin: (Wallac CR28-100)

Tris Hydrochloride

96 well plates: polypropylene for intermediate dilutions (Costar #3794) and either a clear-bottomed white SPA plates (Costar #3632) or a black

15 Polyfiltronics plate (UP350 PSB) for assays.

RESULTS

Ligands, in particular CDCA and the corresponding glycine and taurine conjugates, increased the interaction between FXR and the SRC1 peptide as determined with time-resolved FRET (Fig. 3A). Dose response analysis showed that CDCA, glyco-CDCA and tauro-CDCA increased the amount of SRC1 peptide bound to the FXR LBD. A typical saturable concentration response curve characteristic of specific interaction was observed.

Table 1. Potency of bile acids for binding to FXR as determined in the cell-free ligand sensing assay. The indicated values for CDCA and its conjugates are EC₅₀s derived from dose response analysis as described in Fig 3B. The indicated values for other bile acids and their conjugates are half-maximal inhibitor concentrations IC₅₀s derived from dose response analysis as described in Fig 3C.

30

Bile Acid	free acid	glycine- conjugate	taurine- conjugate
-----------	--------------	-----------------------	-----------------------

CDCA	4.5 μ M	10 μ M	10 μ M
CA	>1000 μ M	>1000 μ M	>1000 μ M
LCA	3.8 μ M	4.7 μ M	3.8 μ M
DCA	100 μ M	\geq 500 μ M	\geq 500 μ M
MCA	>1000 μ M	Not tested	\geq 500 μ M

Although LCA, DCA, and (*E*)-[(tetrahydrotetramethylnaphthalenyl)-propenyl]-benzoic acid (TTNPB) activate FXR in the cell-based reporter assay (Fig. 2B), they did not promote interactions between the FXR LBD and SRC1 in the FRET ligand-sensing assay (Fig. 3A). However, when these compounds were assayed in the presence of 50 μ M CDCA, they disrupted the CDCA-FXR-SRC1 complex in a dose-dependent fashion (Fig. 3C and Table 1). Thus, CDCA, LCA, DCA, and TTNPB are ligands for FXR. Similarly, the taurine and glycine conjugates of LCA decreased the fluorescence signal, indicating displacement of CDCA from FXR with IC₅₀ values of 3.8 and 4.7 μ M, respectively (Table 1). The conjugated forms of DCA and MCA caused a small decrease in fluorescence at the highest concentration tested, and no effect was obtained with the conjugated forms of CA and ursodeoxycholic acid (Table 1).

As used in the claims for this assay, "nuclear receptor coactivator peptide" means a peptide whose affinity for the receptor is changed in the presence of ligand and which has a LXXLL (SEQ ID NO.:6) motif. Examples of coactivator peptides useful for ligand identification by this method that have been demonstrated to interact with FXR include SRC-1 and those listed below:

1. B-QEQLSPKKKENNALLRYLLDRDDPS-CONH₂ (SEQ ID NO.: 2), ACTR (734-758), RAC3 (724-748), SRC-3 (724-748), AIB1 (724-748), pCIP (716-740)
2. B-QEPVSPKKKENALLRYLLDKDDTKD-CONH₂ (SEQ ID NO.:3), TIF2 (732-756)
3. B-GSTHGTSLKEKHKILHRLQLDSSSPVD-CONH₂ (SEQ ID NO.:4), TIF2 (676-702)
- 25 4. B-SNMHGSLLQEKRILHKLLQNGNSPAE-CONH₂ (SEQ ID NO.:5), pCIP (664-690), RAC3 (671-697), ACTR (681-707), AIB1 (671-697)

The sequences of peptides 1 and 4 appear in a number of coactivators, hence the multiple names. The "B" in the sequences stands for biotinylated, which is a modification that allows attachment of the peptide during the analysis.

5

Example 5 The present invention provides the non-steroidal compound GW4064 as a potent and selective FXR ligand.

Synthesis of GW4064

A solution of 2,6-dichlorobenzaldehyde (25g, 0.14 mole) in ethanol (200 mL) was added to a solution of hydroxylamine hydrochloride (11g, 0.16 mole) and sodium hydroxide (6.3g, 0.16 mole) in water (100 mL). The resulting mixture was stirred at 90°C for 24 hours. The volume was reduced in vacuo by ca 30 mL which induced a precipitate. The flask was then cooled to room temperature and the white solids were collected by filtration and washed with water (2x 100 mL). Yield = 25.9g. (96%) of 2,6-dichlorobenzaldehyde oxime. A 500 mL round bottom flask was charged with a solution of 2,6-dichlorobenzaldehyde oxime (13g, 0.07 mole) in N,N-dimethyl formamide (150 mL). The flask was placed in an ambient temperature water bath. The flask was then charged with N-chlorosuccinimide (9.2g, 0.07 mole). Within minutes of dissolution, an exotherm was observed along with a significant color change to dark yellow. The reaction was stirred an additional hour then the contents were then poured into water (200 mL) and the product extracted with diethyl ether (300mL). The ethereal layer was washed with water (3x 100 mL) and brine (50mL), then dried over anhydrous magnesium sulfate. After filtering, the solvent was removed in vacuo to yield 14.5g of a yellow oil. (94%) of 2,6-dichlorophenyl hydroximic chloride which was used without further purification. A stirred solution of methyl isobutyryl acetate (2g, 15.6 mmol) in tetrahydrofuran (15 mL) was treated with a solution of sodium methoxide (31.5 mL, 0.5 M in methanol) followed by a solution of 2,6-dichlorophenyl hydroximic chloride (3.5g, 15.6 mmol) in tetrahydrofuran (5 mL). After stirring at ambient temp 16h the solvent was removed in vacuo. The resulting residue was

partitioned with diethyl ether (100 mL) and water (100 mL). The ethereal layer was washed with brine (50 mL), dried over anhydrous magnesium sulfate, filtered and condensed to an oil. The product was purified by flash chromatography on silica gel using 10% ethyl acetate in hexane as mobile phase. Yield = 3.1 g. (62%) of 3-(2,6-dichlorophenyl)-4-carbomethoxy-5-isopropyl-isoxazole.

A solution of 3-(2,6-dichlorophenyl)-4-carbomethoxy-5-isopropyl-isoxazole (8.4 g, 27 mmol) in tetrahydrofuran (60 mL) was cooled to 0°C under a nitrogen atmosphere while a solution of diisobutylaluminum hydride (38 mL, 10 1.5M in toluene) was added dropwise. The reaction was allowed to warm slowly to ambient temperature over several hours and stir at room temperature 16h total. The flask was again cooled to 0°C and methanol (2 mL) was carefully added over a 10 minute period. Water (20 mL) was added dropwise and a gelatinous mixture formed. Sodium hydroxide (30mL, 2N) was added 15 and the material was filtered over a plug of celite. After the solids were nearly dry, they were extracted with ethyl acetate (4x 50 mL) and the filtrates were combined. The organic layer was washed with water (2x 50 mL), brine (50 mL), were dried over anhydrous magnesium sulfate, filtered and condensed to a white crystalline solid. Yield = 7.1 g. (93%) of 3-(2,6-dichlorophenyl)-4-hydroxymethyl-5-isopropyl-isoxazole.

20 A solution of 2-chloro-4-hydroxy-benzaldehyde (0.5 g, 3.2 mmol) and 3-(2,6-dichlorophenyl)-4-hydroxymethyl-5-isopropyl-isoxazole (0.9 g, 3.2 mmol) and triphenylphosphine (0.84 g, 3.2 mmol) in dichloromethane (10 mL) was treated with diisopropyl azodicarboxylate (0.63 mL, 3.2 mmol) dropwise. A brief 25 exotherm was observed and the reaction was stirred at ambient temperature 4 hours. The solvent was removed in vacuo and the resulting residue was purified by flash chromatography on silica gel using dichloromethane as the mobile phase. Yield = 0.86 g. (63%) of 3-(2,6-dichlorophenyl)-4-(3'-chloro-4'-formylphenoxy)-methyl-5-isopropyl-isoxazole.

30 Methyl-3-bromomethyl-benzoate (14 g, 61 mmol) in triethylphosphite (12 mL, 70 mmol) was stirred at 185°C for 4 hours. Excess triethylphosphite was

removed by vacuum distillation (150 mT, 145° C) and the resulting oil was used without further purification. Yield = 17.5 g. (100%) of diethyl (3-methoxycarbonyl) benzyl phosphonate.

A solution of diethyl (3-methoxycarbonyl) benzyl phosphonate (0.68 g, 2.4 mmol) in tetrahydrofuran (5 mL) was treated with sodium hydride (0.1 g, 2.5 mmol 60% dispersion in oil). Vigorous gas evolution was observed. Once the bubbling subsided, 3-(2,6-dichlorophenyl)-4-(3'-chloro-4'-formylphenoxy)-methyl-5-isopropyl-isoxazole (0.85 g, 2 mmol) as a solution in tetrahydrofuran (3 mL) was added dropwise and the reaction was stirred 5 hours at ambient temperature. The volume was diluted to 70 mL with ethyl acetate and the mixture was washed with water (50 mL), saturated sodium bisulfate (20 mL), and brine (50 mL). The organic phase was dried over anhydrous magnesium sulfate, filtered, and condensed to an oil. This material was purified by flash chromatography on silica gel using dichloromethane as the mobile phase. Yield = 0.95 g. (86%) of 3-(2,6-dichlorophenyl)-4-(3'-carbomethoxy-2-chloro-stilben-4-yl)-oxymethyl-5-isopropyl-isoxazole.

A solution of (6.3 g, 11.3 mmol) in tetrahydrofuran (25 mL) was treated with lithium hydroxide (25 mL, 1M in water) and the mixture was stirred vigorously at room temperature 24 hours. The tetrahydrofuran was removed in vacuo and the resulting residue was partitioned with ethyl acetate (300 mL), water (100 mL), and saturated sodium bisulfate (5 mL). The organic phase was washed with water (50 mL), brine (50 mL) then was dried over anhydrous magnesium sulfate, filtered, and condensed to yield a yellow residue. The product was purified by flash chromatography on silica gel using ethyl acetate as the mobile phase. Yield = 6.0 g, (98%) of 3-(2,6-dichlorophenyl)-4-(3'-carboxy-2-chloro-stilben-4-yl)-oxymethyl-5-isopropyl-isoxazole.

GW4064 activates rodent FXR with a half maximal effective concentration (EC_{50}) of approximately 300 nM in cell-based reporter assays. Tissues have been examined from vehicle or GW4064-treated rats for genes that are differentially expressed. FXR target genes have been identified in the liver and intestine, two tissues where FXR is abundantly expressed. Our data provide

evidence for a broad role for FXR in lipid homeostasis and digestion.

Furthermore, our data indicate utilities for FXR ligands in the treatment of diseases associated with impaired/altered lipid homeostasis and digestion.

Fischer rats were treated with either vehicle alone or 30 mg/kg of the 5 FXR-selective ligand GW4064 for 7 days. At the end of the study, serum lipid levels were determined. Serum triglyceride levels were decreased by 26% in GW4064-treated animals. Similarly, natural bile acids, which serve as FXR ligands, lower triglyceride levels when administered to rodents or man (Iser and Sali, 1981). Our data with the non-steroidal FXR ligand GW4064 demonstrate 10 that FXR mediates the triglyceride-lowering effects of bile acids.

Liver and small intestine were removed from GW4064- and vehicle-treated Fischer rats and cDNA prepared. Curagen GeneCall technology was employed to identify genes that were differentially regulated by GW4064 in these tissues.

15

Intestine

Nearly twenty genes were identified in the intestine that were regulated >1.5-fold by GW4064 (Table 2). The expression of roughly half of these genes was decreased by GW4064 treatment. Notably, all of these downregulated genes 20 are involved in either lipid absorption or proteolysis. Among the genes involved in lipid absorption are three lipases (lipase, pancreatic lipase, and pancreatic phospholipase A2), a colipase, and the intestinal fatty acid binding protein. The proteases carboxypeptidase A1, carboxypeptidase A2, carboxypeptidase B, pancreatic trypsin I, cationic trysinogen, pancreatic trysinogen II, elastasel, 25 and elastase II were all down regulated ≥ 2 fold. Finally, pancreatic amylase, which hydrolyzes starches and glycogen, was downregulated >8 -fold. Taken together, these data provide evidence that FXR plays an important role in the coordinate regulation of digestion in the small intestine, including the metabolism of fatty acids, protein, and carbohydrates. Moreover, these data 30 suggest that bile acids, as natural FXR ligands, serve a broader role in the regulation of digestion in the small intestine than was previously appreciated.

Among the genes whose expression was increased by GW4064 in the intestine was the previously established FXR target gene IBABP. Expression of hepatic squalene synthetase, an enzyme essential for cholesterol biosynthesis, and the orphan nuclear receptor short heterodimerizing partner 1 (SHP-1) was also increased in the intestine. SHP-1 may be involved in the coordinate down-regulation of genes modulated by nuclear receptors, as discussed below.

Table 2

A. Genes Downregulated in Liver

- 10 apolipoprotein B -2.7
plasma proteinase inhibitor alpha-1-inhibitor III group 3 (m22360) -2.7
L-gulono-gamma-lactone oxidase (d12754) -2.2
Peroxisomal enoyl-CoA:hydratase-3-hydroxyacyl-CoA bifunctional enzyme (k03249) -2.0
15 liver fatty acid binding protein (L-FABP, m13501) -2.0
CYP4A2 (m57719) -1.9
CYP3A23 (x96721) -1.8
CYP3A1 (x64401) -1.6

20 **B. Genes Upregulated in Liver**

- Small heterodimer partner homolog (d86580) +6.6
insulin-induced growth-response protein, CL-6 (I13619) +5.9
elongation factor 2, EF-2 (y07504) +4.3
mouse cornichon +4.0
25 protein kinase C receptor (u03390) +3.1
mitochondrial cytochrome c oxidase (m27315) +2.7
cystathione gamma-lyase (x53460, d17370) +2.3
cytosolic phosphoenolpyruvate carboxykinase (k03243) +2.1
histidase (m58308) +2.1
30 S-adenosylmethionine synthetase (x60822) +2.0
lanosterol 14-alpha-demethylase (u17697)+1.9

G protein-coupled purinoceptor P2U (I46865) +1.9
hepatic squalene synthetase (m95591) +1.5

5 Liver

Unlike the intestine, most liver genes whose expression was affected by GW4064 were upregulated (Table 3). Expression of the gene encoding the orphan nuclear receptor SHP-1 was increased the most, nearly 7-fold. SHP-1 was originally cloned in a yeast two hybrid assay based on its interaction with 10 the orphan nuclear receptor CAR (Seol et al., 1996). Unlike most nuclear receptors, SHP-1 lacks a DNA-binding domain. In addition to CAR, SHP-1 interacts with other nuclear receptors including thyroid hormone receptor, retinoic acid receptor, RXR, estrogen receptor α , estrogen receptor β , peroxisome proliferator-activated receptor α , and hepatocyte nuclear receptor 4 15 α . In cell-based reporter assays, SHP-1 specifically inhibits transactivation by the family members with which it interacts (Johansson et al., 1999; Masuda et al., 1997; Seol et al., 1996). Thus, the induction of SHP-1 gene expression by FXR ligands may, in turn, result in decreases in expression of genes regulated by nuclear receptors. This mechanism is likely to have important implications 20 in the regulation of CYP7a gene expression by bile acids (see below).

Table 3

Genes Having Altered Expression in Intestine:

lipase (x61925) -9.5
pancreatic lipase (d88534) -4.4
25 colipase (m58370) -10.2
pancreatic phospholipase A-2 (d00036) -7.4
pancreatic amylase (m24962) -8.1
carboxypeptidase A1 (m23986) -6.1
carboxypeptidase A2 (m23721) -5.4
30 carboxypeptidase B (m23959) -4.2

pancreatic trypsin I (j00778) -6.8
pancreatic cationic trypsinogen (m16624)
pancreatic trypsinogen II (v01274) -3.6
elastase I (v01234, l00112) -2.0
5 elastase II (l00118, l00124) -4.6
I-BABP (l22788) +2.3
intestinal fatty acid binding protein (FABP, k01180) -1.8
hepatic squalene synthetase (m95591) +1.9
protein kinase C receptor (u03390) +1.6
10 elongation factor 2, EF-2 (y07504) +2.4
Small heterodimer partner homolog (d86580) +2.2

GW4064 treatment induced the expression of two enzymes required for de novo cholesterol biosynthesis, lanosterol 14-alpha demethylase and 15 squalene synthetase.

GW4064 treatment increased the expression of a number of genes whose products have well-established roles in various cellular processes including amino acid metabolism (cystathionase, histidase, S-adenosylmethionine), glucose biosynthesis (phosphoenolpyruvate 20 carboxykinase), protein translation (elongation factor 2), and electron transport (mitochondrial cytochrome c oxidase). GW4064 treatment also increased expression of the activated protein kinase C binding protein RACK1, the P2U ATP receptor, cornichon, and insulin-induced growth-response protein CL-6. The latter gene is interesting in that it is abundantly expressed in proliferating 25 hepatocytes during development and regeneration.

Consistent with the observed down-regulation of genes involved in fatty acid absorption in the intestine, GW4064 treatment decreased expression of genes involved in hepatic fatty acid metabolism, including fatty acid oxidation (peroxisomal bifunctional protein, CYP4A2), transport (liver fatty acid binding 30 protein), and secretion (apolipoprotein B). Thus FXR appears have a coordinate role in the regulation of fatty acid homeostasis in the intestine and

liver. GW4064 treatment also decreased expression of genes involved in steroid and xenobiotic metabolism (CYP3A23), vitamin C biosynthesis (L-gulono-gamma-lactone oxidase), and inflammatory response (plasma proteinase inhibitor alpha-1-inhibitor group III).

5 Results

Our results implicate FXR, and by inference its bile acid ligands, in the coordinate regulation of genes involved in lipid absorption and digestion in the small intestine and lipid homeostasis in the liver. Moreover, our results suggest several mechanisms whereby FXR may regulate CYP7a expression in
10 the liver.

Lipid absorption and digestion.

Our data indicate that FXR regulates a program of genes involved in digestive processes in the small intestine. Thus, a model for FXR effects in the small
15 intestine has been drawn. The ingestion of dietary fatty acids results in cholecystokinin (CCK) release from specialized endocrine cells in the small intestine. CCK, in turn, effects the release of proteases required for digestion and the contraction of the gall bladder, which releases bile acids into the duodenum. The released bile acids may activate FXR and result in decreases
20 in the expression of lipases, a colipase, and proteases in the small intestine. Thus, FXR may serve as a central regulator in the feedback regulation of digestion, sensing bile acids released during the course of digestion and reducing the expression of genes involved in the digestion and absorption of fatty acids and protein. It is well established that increases in bile acids reduce
25 triglyceride levels in mammals (Iser and Sali, 1981). The present model suggests that these decreases in triglyceride levels are due in part to decreased absorption of fatty acids in the small intestine.

The present data also indicate that FXR has coordinate effects on the regulation of fatty acid metabolism in the liver. Consistent with decreased lipid
30 absorption in the intestine, the expression levels of genes involved in hepatic oxidation, transport (liver fatty acid binding protein), and secretion (apoB) of

fatty acids are decreased by an FXR ligand.

Implications of FXR in the regulation of CYP7a

Two possible mechanisms for bile acid feed back and regulate the expression
5 of CYP7a can thus be drawn. First, expression of the orphan receptor SHP-1 was dramatically increased in the livers of GW4064-treated rats. SHP-1 is known to heterodimerize with and suppress the activity of a number of nuclear receptors, including thyroid hormone receptor and hepatocyte nuclear receptor 4- α , both of which have been implicated in the regulation of CYP7a expression
10 (Johansson et al., 1999; Masuda et al., 1997; Seol et al., 1996). The present invention provides that SHP-1 expression was also increased in the small intestine of GW4064-treated rats. FXR-mediated increases in SHP-1 gene expression may account in part for the down-regulation of gene expression that is observed in both the liver and small intestine of GW4064-treated rats. An
15 interesting extension is that compounds that bind to SHP-1 and block its ability to antagonize other nuclear receptors may have therapeutic utility in the up-regulation of CYP7a expression and increased metabolism of cholesterol to bile acids.

GW4064 treatment also resulted in increases in the expression of
20 RACK1, which binds activated protein kinase C. RACK1 is a shuttling protein that moves activated protein kinase C between intracellular sites (Rodriguez et al., 1999). Interestingly, the activation of protein kinase C has been shown to repress CYP7a expression in primary cultures of rat hepatocytes and to inhibit bile flow in the perfused rat liver model (Bouscarel et al., 1999). Thus, FXR-mediated effects on protein kinase C signaling pathways may impact the expression of CYP7a and perhaps other genes.

The compounds of Formula (I) or (II) can be synthesized using standard techniques of organic chemistry. A convergent strategy can be employed in which a hydroxystilbene and a hydroxymethyisoxazole are prepared
30 independently and then condensed using a Mitsunobu coupling to generate the ether linkage. Compounds with anilino linkages can be prepared by converting

the hydroxyl residue of a hydroxymethyisoxazole into a leaving group, such as bromide or mesylate, followed by reaction with an aminostilbenes.

Hydroxymethylsoxazoles can be prepared by the condensation of a β -keto ester enolates with an α -halo-substituted hydroximic acid. The resulting 5 esters can be reduced to an alcohol with a metal hydride reducing agent such as DIBAL.

Hydroxystilbenes can be prepared by Horner-Wadsworth-Emmons coupling of an aryl aldehyde and an arylmethylene phosphonate ester, or by Heck coupling of a styrene with an arylbromide, iodide, or triflate in the 10 presence of a palladium catalyst. Using standard chemical methods, tritium or iodine 125 can be incorporated into the compounds of formula (I) and (II).

The compounds of Formula (I) or (II) and their pharmaceutically acceptable derivatives are conveniently administered in the form of pharmaceutical compositions. Thus, in another aspect of the invention, we 15 provide a pharmaceutical composition comprising a compound of formula (I) or (II) or a pharmaceutically acceptable derivative thereof adapted for use in human or veterinary medicine. Such compositions may conveniently be presented for use in conventional manner in admixture with one or more physiologically acceptable carriers or excipients.

20 The compounds of Formula (I) or (II) and their pharmaceutically acceptable derivatives may be formulated for administration in any suitable manner. They may, for example, be formulated for topical administration or administration by inhalation or, more preferably, for oral, transdermal or parenteral administration. The pharmaceutical composition may be in a form 25 such that it can effect controlled release of the compounds of Formula (I) and (II) and their pharmaceutically acceptable derivatives. A particularly preferred method of administration, and corresponding formulation, is oral administration.

For oral administration, the pharmaceutical composition may take the form of, and be administered as, for example, tablets (including sub-lingual 30 tablets) and capsules (each including timed release and sustained release formulations), pills, powders, granules, elixirs, tinctures, emulsions, solutions,

syrups or suspensions prepared by conventional means with acceptable excipients.

For instance, for oral administration in the form of a tablet or capsule, the active drug component can be combined with an oral, non-toxic

5 pharmaceutically acceptable inert carrier such as ethanol, glycerol, water and the like. Powders are prepared by comminuting the compound to a suitable fine size and mixing with a similarly comminuted pharmaceutical carrier such as an edible carbohydrate, as, for example, starch or mannitol. Flavoring, preservative, dispersing and coloring agent can also be present.

10 Capsules can be made by preparing a powder mixture as described above, and filling formed gelatin sheaths. Glidants and lubricants such as colloidal silica, talc, magnesium stearate, calcium stearate or solid polyethylene glycol can be added to the powder mixture before the filling operation. A disintegrating or solubilizing agent such as agar-agar, calcium carbonate or
15 sodium carbonate can also be added to improve the availability of the medicament when the capsule is ingested.

Moreover, when desired or necessary, suitable binders, lubricants, disintegrating agents and coloring agents can also be incorporated into the mixture. Suitable binders include starch, gelatin, natural sugars such as
20 glucose or beta-lactose, corn sweeteners, natural and synthetic gums such as acacia, tragacanth or sodium alginate, carboxymethylcellulose, polyethylene glycol, waxes and the like. Lubricants used in these dosage forms include sodium oleate, sodium stearate, magnesium stearate, sodium benzoate, sodium acetate, sodium chloride and the like. Disintegrators include, without
25 limitation, starch, methyl cellulose, agar, bentonite, xanthan gum and the like. Tablets are formulated, for example, by preparing a powder mixture, granulating or slugging, adding a lubricant and disintegrant and pressing into tablets. A powder mixture is prepared by mixing the compound, suitably comminuted, with a diluent or base as described above, and optionally, with a
30 binder such as carboxymethylcellulose, an aliginate, gelatin, or polyvinyl pyrrolidone, a solution retardant such as paraffin, a resorption accelerator such

as a quaternary salt and/or an absorption agent such as bentonite, kaolin or dicalcium phosphate. The powder mixture can be granulated by wetting with a binder such as syrup, starch paste, acadia mucilage or solutions of cellulosic or polymeric materials and forcing through a screen. As an alternative to
5 granulating, the powder mixture can be run through the tablet machine and the result is imperfectly formed slugs broken into granules. The granules can be lubricated to prevent sticking to the tablet forming dies by means of the addition of stearic acid, a stearate salt, talc or mineral oil. The lubricated mixture is then compressed into tablets. The compounds of the present
10 invention can also be combined with free flowing inert carrier and compressed into tablets directly without going through the granulating or slugging steps. A clear or opaque protective coating consisting of a sealing coat of shellac, a coating of sugar or polymeric material and a polish coating of wax can be provided. Dyestuffs can be added to these coatings to distinguish different unit
15 dosages.

Oral fluids such as solution, syrups and elixirs can be prepared in dosage unit form so that a given quantity contains a predetermined amount of the compound. Syrups can be prepared by dissolving the compound in a suitably flavored aqueous solution, while elixirs are prepared through the use of
20 a non-toxic alcoholic vehicle. Suspensions can be formulated by dispersing the compound in a non-toxic vehicle. Solubilizers and emulsifiers such as ethoxylated isostearyl alcohols and polyoxyethylene sorbitol ethers, preservatives, flavor additive such as peppermint oil or saccharin, and the like can also be added.

25 Where appropriate, dosage unit formulations for oral administration can be microencapsulated. The formulation can also be prepared to prolong or sustain the release as for example by coating or embedding particulate material in polymers, wax or the like.

The compounds of the present invention can also be administered in the
30 form of liposome delivery systems, such as small unilamellar vesicles, large unilamellar vesicles and multilamellar vesicles. Liposomes can be formed from

a variety of phospholipids, such as cholesterol, stearylamine or phosphatidylcholines.

The compounds of the present invention can also be administered in the form of liposome emulsion delivery systems, such as small unilamellar vesicles, large unilamellar vesicles and multilamellar vesicles. Liposomes can be formed from a variety of phospholipids, such as cholesterol, stearylamine or phosphatidylcholines.

Compounds of the present invention may also be delivered by the use of monoclonal antibodies as individual carriers to which the compound molecules are coupled. The compounds of the present invention may also be coupled with soluble polymers as targetable drug carriers. Such polymers can include polyvinylpyrrolidone, pyran copolymer, polyhydroxypropylmethacrylamide-phenol, polyhydroxyethylaspartamidephenol, or polyethyleneoxidepolylysine substituted with palmitoyl residues. Furthermore, the compounds of the present invention may be coupled to a class of biodegradable polymers useful in achieving controlled release of a drug, for example, polylactic acid, polepsilon caprolactone, polyhydroxy butyric acid, polyorthoesters, polyacetals, polydihydropyrans, polycyanoacrylates and cross-linked or amphipathic block copolymers of hydrogels.

The present invention includes pharmaceutical compositions containing 0.1 to 99.5%, more particularly, 0.5 to 90% of a compound of the formula (II) in combination with a pharmaceutically acceptable carrier.

Likewise, the composition may also be administered in nasal, ophthalmic, otic, rectal, topical, intravenous (both bolus and infusion), intraperitoneal, intraarticular, subcutaneous or intramuscular inhalation or insufflation form, all using forms well known to those of ordinary skill in the pharmaceutical arts.

For transdermal administration, the pharmaceutical composition may be given in the form of a transdermal patch, such as a transdermal iontophoretic patch.

- For parenteral administration, the pharmaceutical composition may be given as an injection or a continuous infusion (e.g. intravenously, intravascularly or subcutaneously). The compositions may take such forms as suspensions, solutions or emulsions in oily or aqueous vehicles and may
- 5 contain formulatory agents such as suspending, stabilizing and/or dispersing agents. For administration by injection these may take the form of a unit dose presentation or as a multidose presentation preferably with an added preservative. Alternatively for parenteral administration the active ingredient may be in powder form for reconstitution with a suitable vehicle.
- 10 The compounds of the invention may also be formulated as a depot preparation. Such long acting formulations may be administered by implantation (for example subcutaneously or intramuscularly) or by intramuscular injection. Thus, for example, the compounds of the invention may be formulated with suitable polymeric or hydrophobic materials (for
- 15 example as an emulsion in an acceptable oil) or ion exchange resins, or as sparingly soluble derivatives, for example, as a sparingly soluble salt.
- Alternatively the composition may be formulated for topical application, for example in the form of ointments, creams, lotions, eye ointments, eye drops, ear drops, mouthwash, impregnated dressings and sutures and
- 20 aerosols, and may contain appropriate conventional additives, including, for example, preservatives, solvents to assist drug penetration, and emollients in ointments and creams. Such topical formulations may also contain compatible conventional carriers, for example cream or ointment bases, and ethanol or oleyl alcohol for lotions. Such carriers may constitute from about 1% to about
- 25 98% by weight of the formulation; more usually they will constitute up to about 80% by weight of the formulation.
- For administration by inhalation the compounds according to the invention are conveniently delivered in the form of an aerosol spray presentation from pressurized packs or a nebulizer, with the use of a suitable
- 30 propellant, e.g. dichlorodifluoromethane, trichlorofluoromethane, dichlorotetrafluoroethane, tetrafluoroethane, heptafluoropropane, carbon

dioxide or other suitable gas. In the case of a pressurized aerosol the dosage unit may be determined by providing a valve to deliver a metered amount. Capsules and cartridges of e.g. gelatin for use in an inhaler or insufflator may be formulated containing a powder mix of a compound of the invention and a 5 suitable powder base such as lactose or starch.

- The pharmaceutical compositions generally are administered in an amount effective for treatment or prophylaxis of a specific condition or conditions. Initial dosing in human is accompanied by clinical monitoring of symptoms, such symptoms for the selected condition. In general, the compositions are
- 10 administered in an amount of active agent of at least about 100 µg/kg body weight. In most cases they will be administered in one or more doses in an amount not in excess of about 20 mg/kg body weight per day. Preferably, in most cases, dose is from about 100 µg/kg to about 5 mg/kg body weight, daily. For administration particularly to mammals, and particularly humans, it is
- 15 expected that the daily dosage level of the active agent will be from 0. 1 mg/kg to 10 mg/kg and typically around 1 mg/kg. It will be appreciated that optimum dosage will be determined by standard methods for each treatment modality and indication, taking into account the indication, its severity, route of administration, complicating conditions and the like. The physician in any event will determine
- 20 the actual dosage which will be most suitable for an individual and will vary with the age, weight and response of the particular individual. The effectiveness of a selected actual dose can readily be determined, for example, by measuring clinical symptoms or standard anti-inflammatory indicia after administration of the selected dose. The above dosages are exemplary of the average case. There
- 25 can, of course, be individual instances where higher or lower dosage ranges are merited, and such are within the scope of this invention. For conditions or disease states as are treated by the present invention, maintaining consistent daily levels in a subject over an extended period of time, e.g., in a maintenance regime, can be particularly beneficial.
- 30 By "therapeutically effective amount" is meant a symptom-alleviating or symptom-reducing amount, a cholesterol-reducing amount, a fatty acid

absorption-blocking amount, a protein and/or carbohydrate digestion-blocking amount and/or a de novo cholesterol biosynthesis -blocking amount of a compound. Such amounts can be readily determined by standard methods, such as by measuring cholesterol levels or observing alleviation of clinical 5 symptoms. For example, the clinician can monitor accepted measurement scores for treatments for these diseases, disorders and conditions.

The abbreviations used in the examples are included below.

	ACTR	Activator for Thyroid Hormone and Retinoid Receptors
10	AIB1	Amplified in Breast Cancer
	APC	Allophycocyanin
	APMSF	p-Amidinophenylmethylsulfonylfluoride, HCl
	Bestatin	[(2S,3R)-3-Amino-2-hydroxy-4-phenylbutanoyl]-L-leucine
	BSA	bovine serum albumin
15	CHAPS	(3-[3-Cholamidopropyl]-dimethylammonio)-1-propanesulfonate
	CPM	counts per minute
	DMSO	dimethylsulfoxide
	DTT	dithiothreitol
	EDTA	Ethylenediaminetetraacetic acid
20	FXR	Farnasoid X Receptor
	IBTG	isopropyl-β-D-thiogalactopyranoside
	LBD	ligand binding domain
	LXR	Liver X Receptor
	OD ₆₀₀	optical density at 600 nm
25	PBS	phosphate buffered saline [100mM NaPhosphate, pH 7.2, 50mM NaCl]
	pCIP	Co-Integrator Protein
	RAC	Receptor Activated Cofactor
	RPM	revolutions per minute
30	RXR	Retinoid X Receptor
	SA-APC	Streptavidin Crosslinked Allophycocyanin

SPA Scintillation Proximity Assay
SRC Steroid Receptor Cofactor
TIF Transcriptional Intermediary Factor
Tris tris-(Hydroxymethyl)-aminomethane

5

The following references are noted and the entire disclosure of each is herein incorporated by reference

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What is claimed is:

1. A method for the rapid determination of a ligand for a Farnesoid X receptor which comprises contacting a component to be tested with an isolated Farnesoid X receptor ligand binding domain which is associated with a first marking component, and a nuclear receptor coactivator peptide associated with a second marking component, and measuring the interaction between the marking components to determine whether the component to be tested modifies binding between the Farnesoid X receptor ligand binding domain and the nuclear receptor coactivator peptide.

10

2. The method of claim 1 wherein the nuclear receptor coactivator peptide is SEQ ID NO.:1, SEQ ID NO.:2, SEQ ID NO.:3, SEQ ID NO.:4, or SEQ ID NO.:5.

15 3. The method of claim 1, wherein the interaction of the marking components is measured by comparing a signal produced by a combination of the nuclear receptor coactivator peptide, the isolated nuclear receptor binding domain and the component to be tested with a signal produced by a combination of the nuclear receptor coactivator peptide and the isolated nuclear receptor ligand binding domain in the absence of the component to be tested.

20
25 4. A method of identifying compounds for the treatment of diseases or disorders modulated by FXR, comprising the step of determining whether the compound interacts directly with FXR, wherein a compound that interacts directly with FXR is a compound for the treatment.

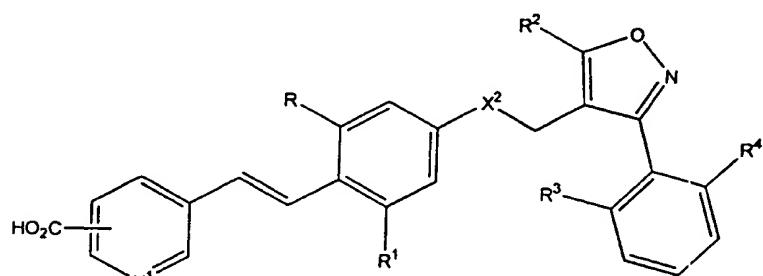
30 5. A method of claim 4 wherein said step of determining comprises the step of specifically detecting the interaction of SRC-1 (LCD2, 677-697) peptide with FXR.

6. A method for the rapid determination of a ligand for Farnesoid X receptor which comprises contacting a component to be tested with an isolated nuclear receptor ligand binding domain which is associated with a first marking component, and a heterodimeric partner for the nuclear receptor ligand binding domain associated with a second marking component, and measuring the interaction between the marking components to determine whether the component to be tested modifies heterodimerization.
- 5

7. The method of claim 6 wherein the heterodimeric partner is an RXR.
- 10

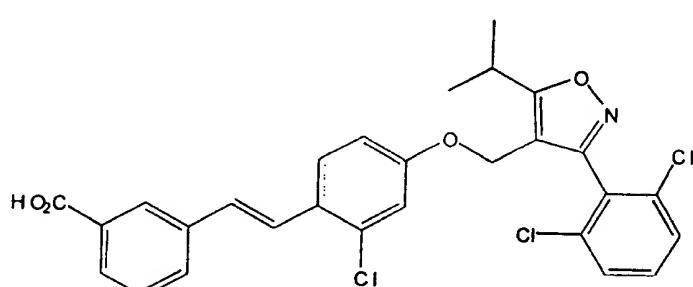
8. A compound identified by the method of any of claims 1-7.

9. A compound that binds Farnesoid X receptor wherein the compound is of the following formula:



- 15
- wherein X¹ is CH, N; X² is O or NH; R and R¹ are independently H, lower alkyl, halogen, or CF₃; R² is lower alkyl; R³ and R⁴ are independently H, lower alkyl, halogen, CF₃, OH, O-alkyl, or O-polyhaloalkyl.

- 20
10. A compound of claim 9 having the following formula:



11. A compound of claim 9 having a detectable label.
12. A method for regulating I-BABP expression in a mammal which comprises activating or inhibiting the Farnesoid X Receptor.
5
13. The method of claim 12, which comprises binding an activating amount of chenodeoxycholic acid to the Farnesoid X Receptor.
14. The method of claim 12, which comprises inhibiting the activation of
10 the Farnesoid X Receptor by chenodeoxycholic acid.
15. A method of regulating the bile transport system in a mammal which comprises activating the Farnesoid X Receptor with a binding ligand.
16. The method of claim 15, wherein the binding ligand is GW4064,
15 chenodexoycholic acid, lithocholic acid, deoxycholic acid, or a glycine or taurine conjugated conjugate derivative thereof.
17. The method of claim 15, wherein the Farnesoid X Receptor is activated
20 by conjugated bile acids in tissues that express bile acid transporters.
18. The method of claim 17, wherein the tissue is in the terminal ileum,
liver, or kidney.
19. The method of claim 15 for regulating intestinal cholesterol absorption,
25 regulating bile acid flux or lowering triglycerides.
20. A method of treating in a mammal a disease which is affected by
cholesterol, triglyceride, or bile acid levels comprising administering to a
30 mammal in need of such treatment a therapeutically effective amount of a ligand for Farnesoid X Receptor.

21. A method of treating atherosclerosis, gallstone disease, lipid disorders, obesity or a cardiovascular disorder, comprising administration of a therapeutically effective amount of a compound which was identified by the
5 method of Claim 4.
22. The method of claim 21 wherein said compound is an activator or inhibitor of FXR heterodimers with the retinoid X receptor.
- 10 23. A method of blocking fatty acid absorption in intestine of a mammal in need of such blocking comprising administering to the mammal a therapeutically effective amount of a Farnesoid X receptor agonist.
- 15 24. The method of claim 23 for treatment of dyslipidemia, obesity or atherosclerosis.
25. A method of blocking protein and carbohydrate digestion in intestine of a mammal in need of such blocking comprising administering to the mammal a therapeutically effective amount of an FXR agonist.
20 ---
26. The method of claim 25 for treatment of obesity.

27. A method of blocking de novo cholesterol synthesis in liver of a mammal in need of such blocking comprising administering to the mammal a therapeutically effective amount of an antagonist of FXR.
25
28. A method of blocking induction of SHP-1 expression in a mammal in need of such blocking comprising administering to the mammal a therapeutically effective amount of an antagonist of FXR.
30
29. A method of blocking SHP-1-mediated repression of CYP7A in a

mammal in need of such blocking comprising administering to the mammal a therapeutically effective amount of an antagonist of SHP-1.

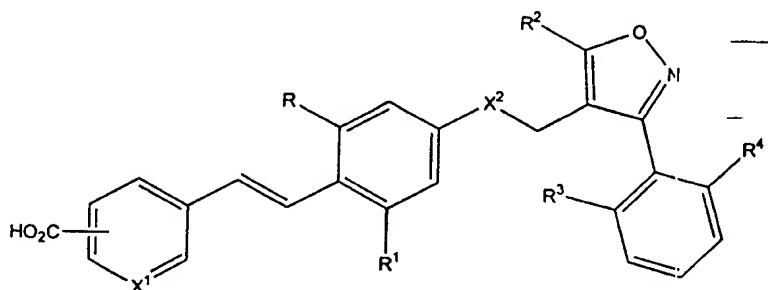
30. The method of any of claims 27-29 for reducing serum cholesterol
5 levels in the mammal.

31. The method of any of claims 27-29 for treatment of atherosclerosis or gall stones.

10 32. The use of an RXR-specific ligand in treatment of a disease or disorder modulated by FXR.

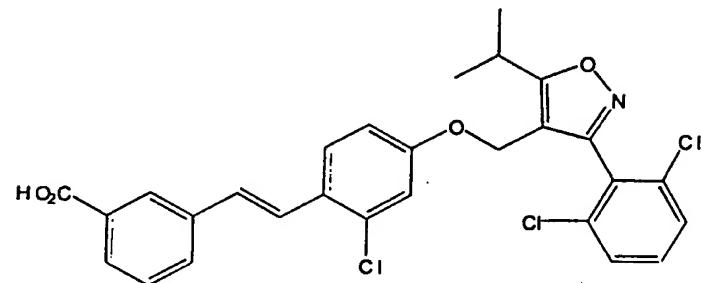
15 33. A method of modulating a gene whose expression is regulated by FXR in a mammal comprising administering to the mammal a therapeutically effective amount of a ligand of FXR.

34. The method of any of claims 15, 20 or 23-33 wherein the ligand is a compound of the formula (I):



35. The method of any of claims 15, 20 or 23-33 wherein the ligand is the following compound (Formula (II)):

5



10

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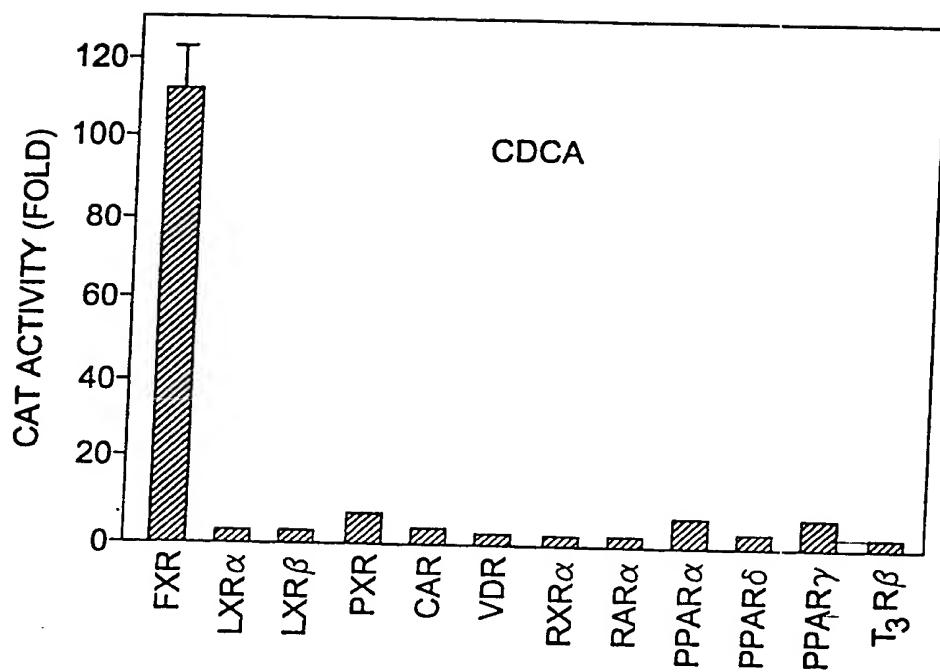
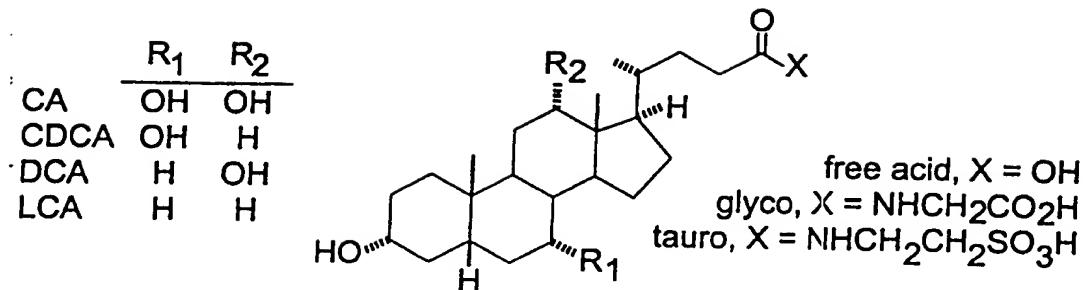
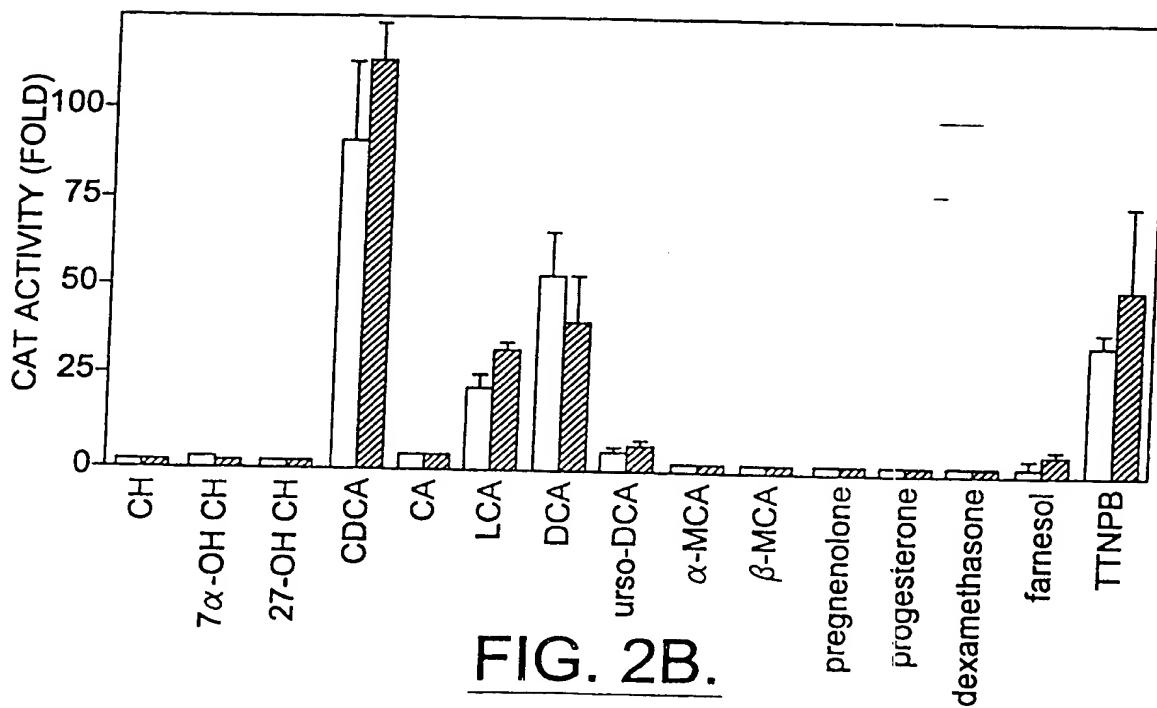
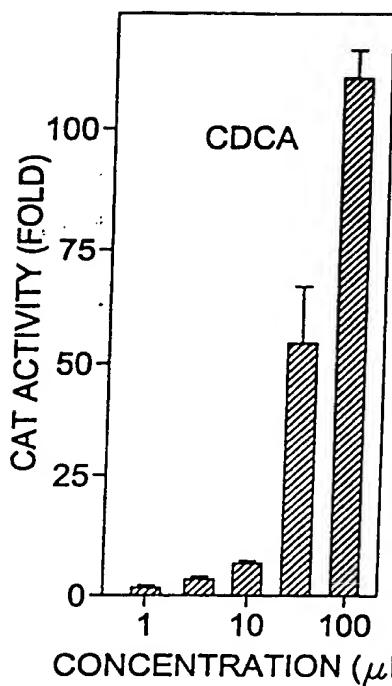
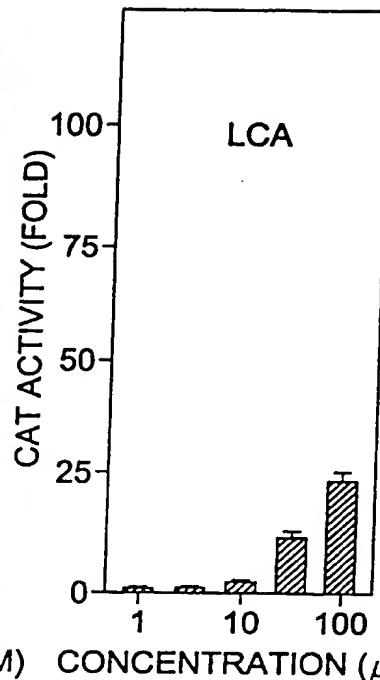
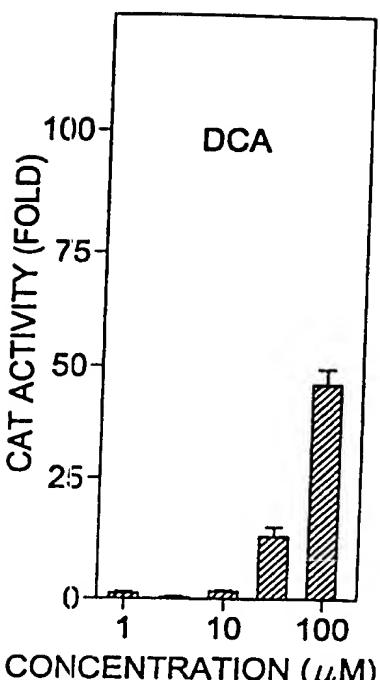
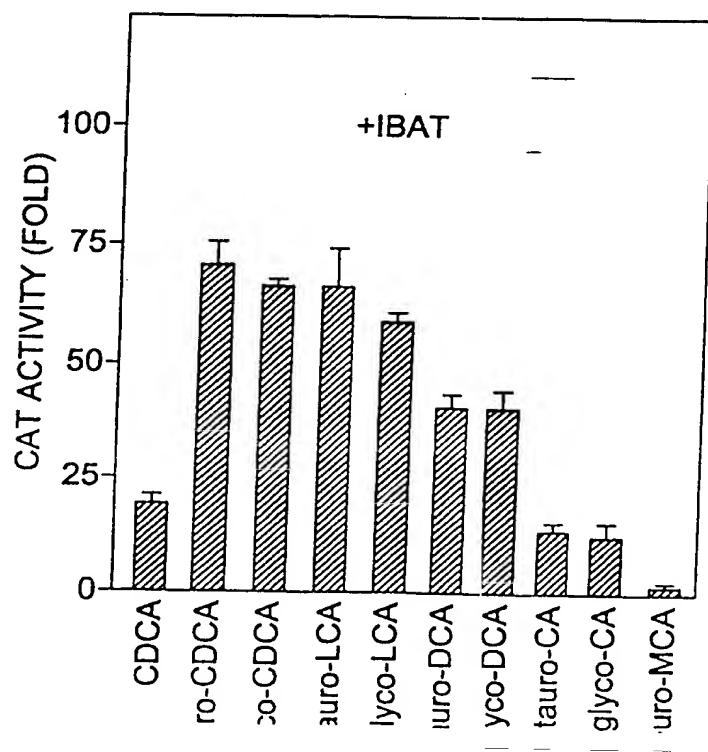
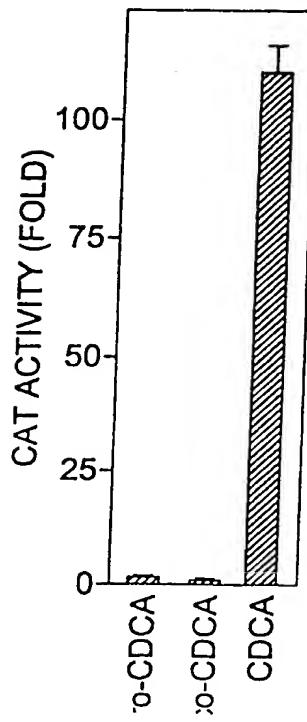


FIG. 1.

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**FIG. 2A.**

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**FIG. 2C.****FIG. 2D.****FIG. 2E.**

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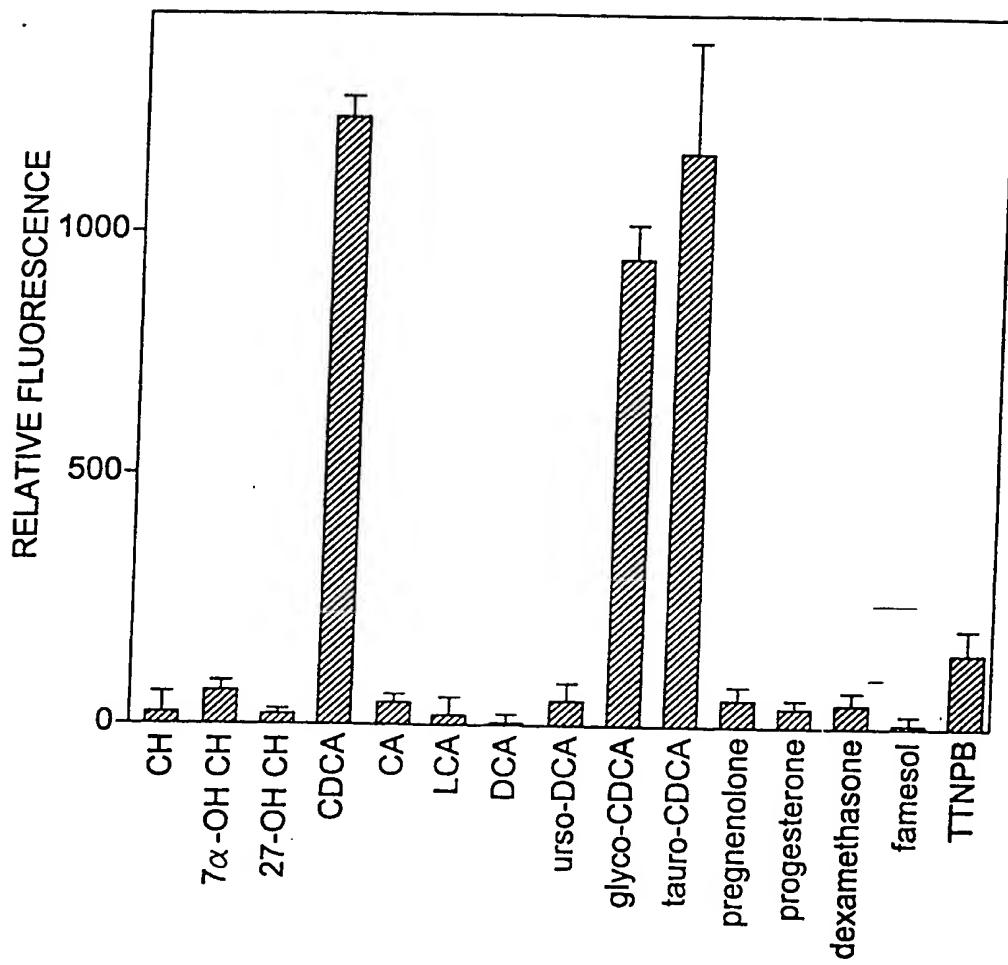


FIG. 3A.

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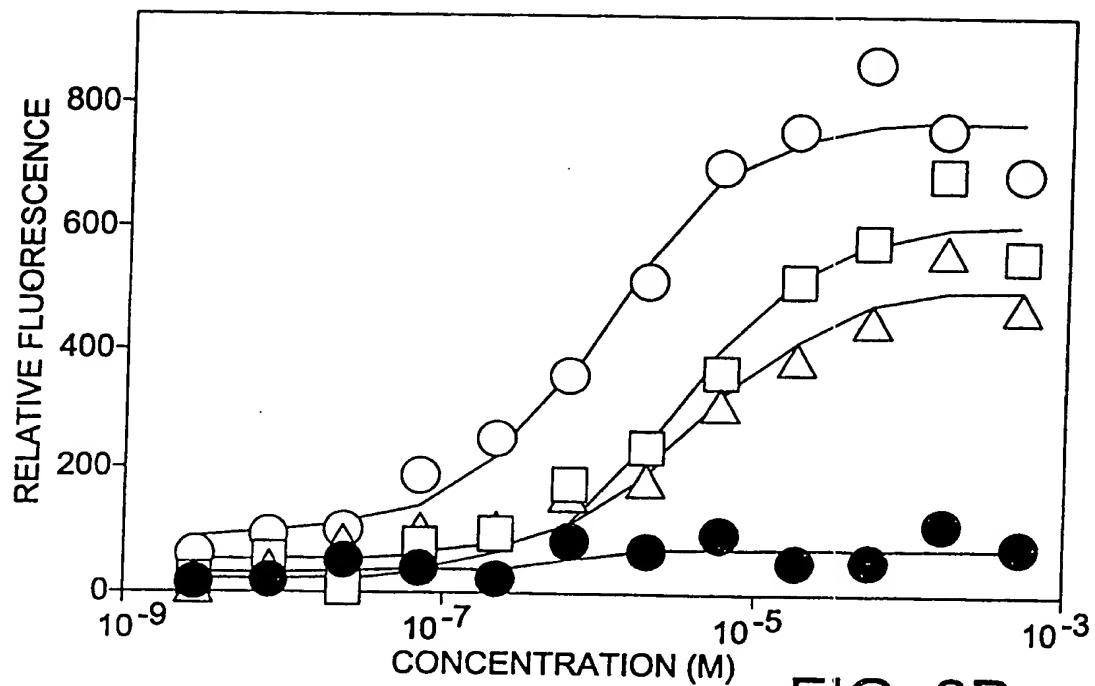
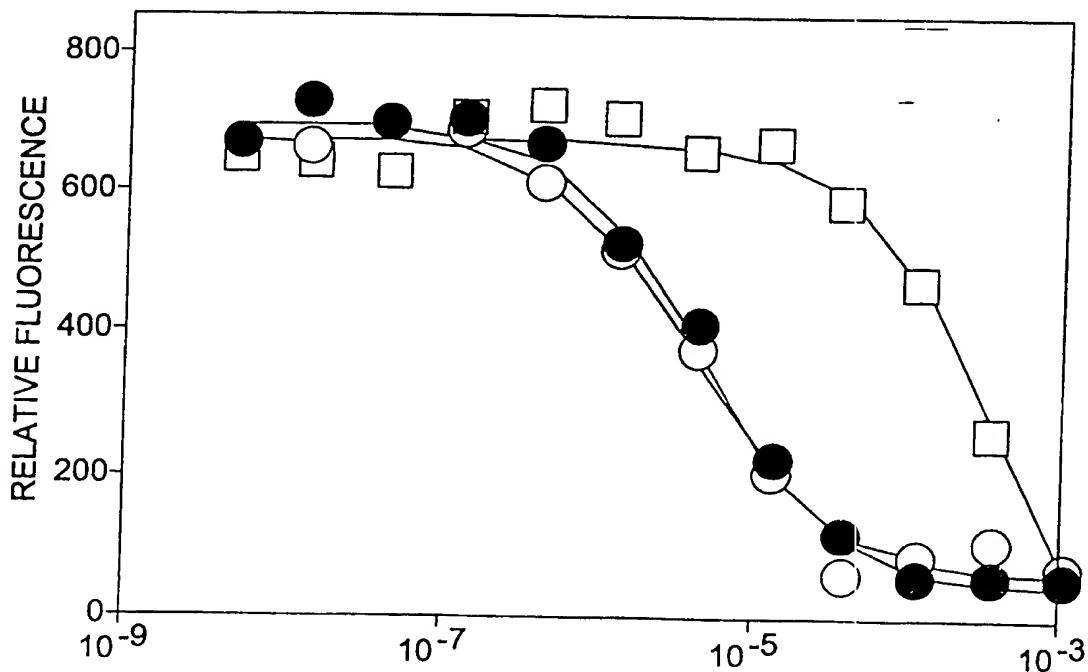
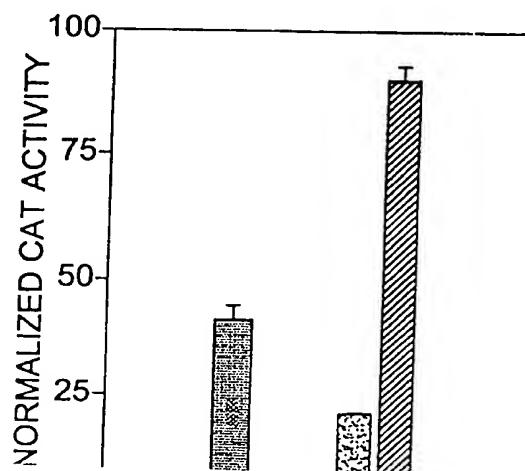
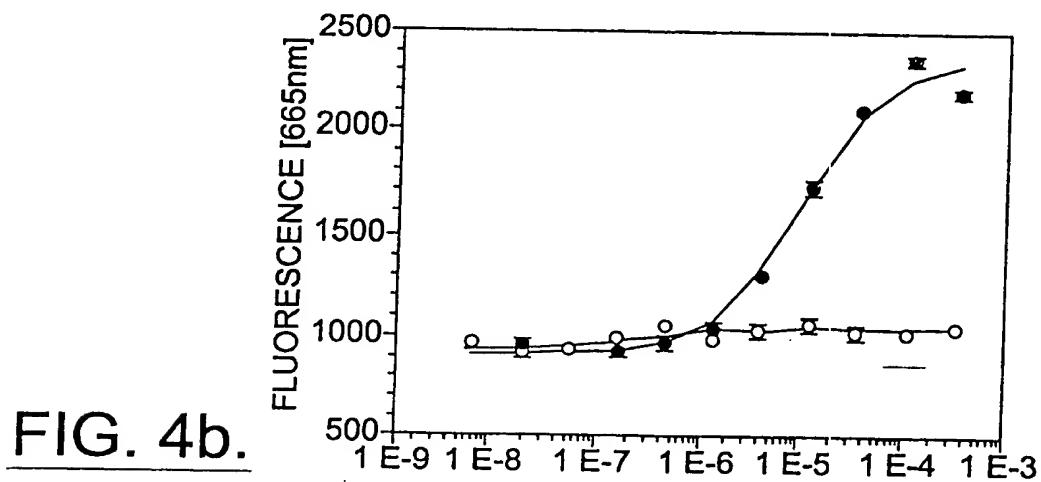
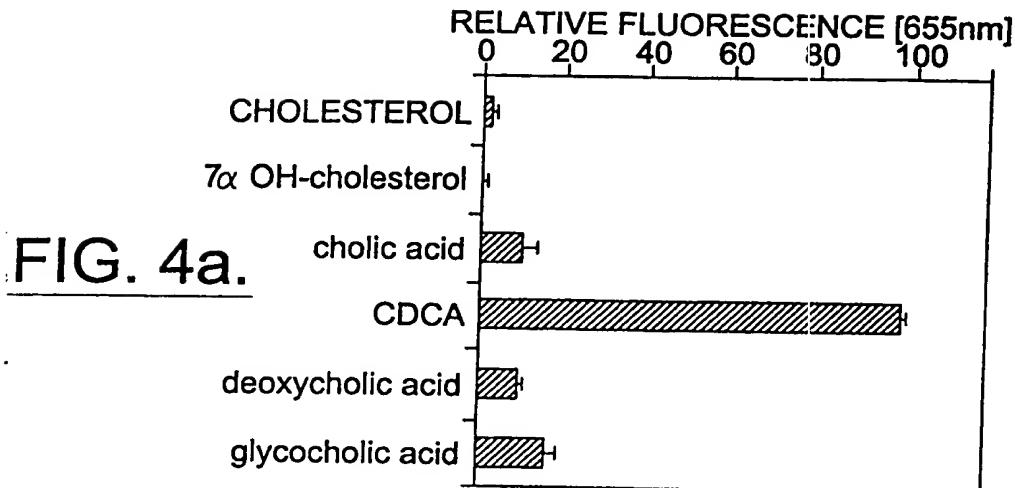


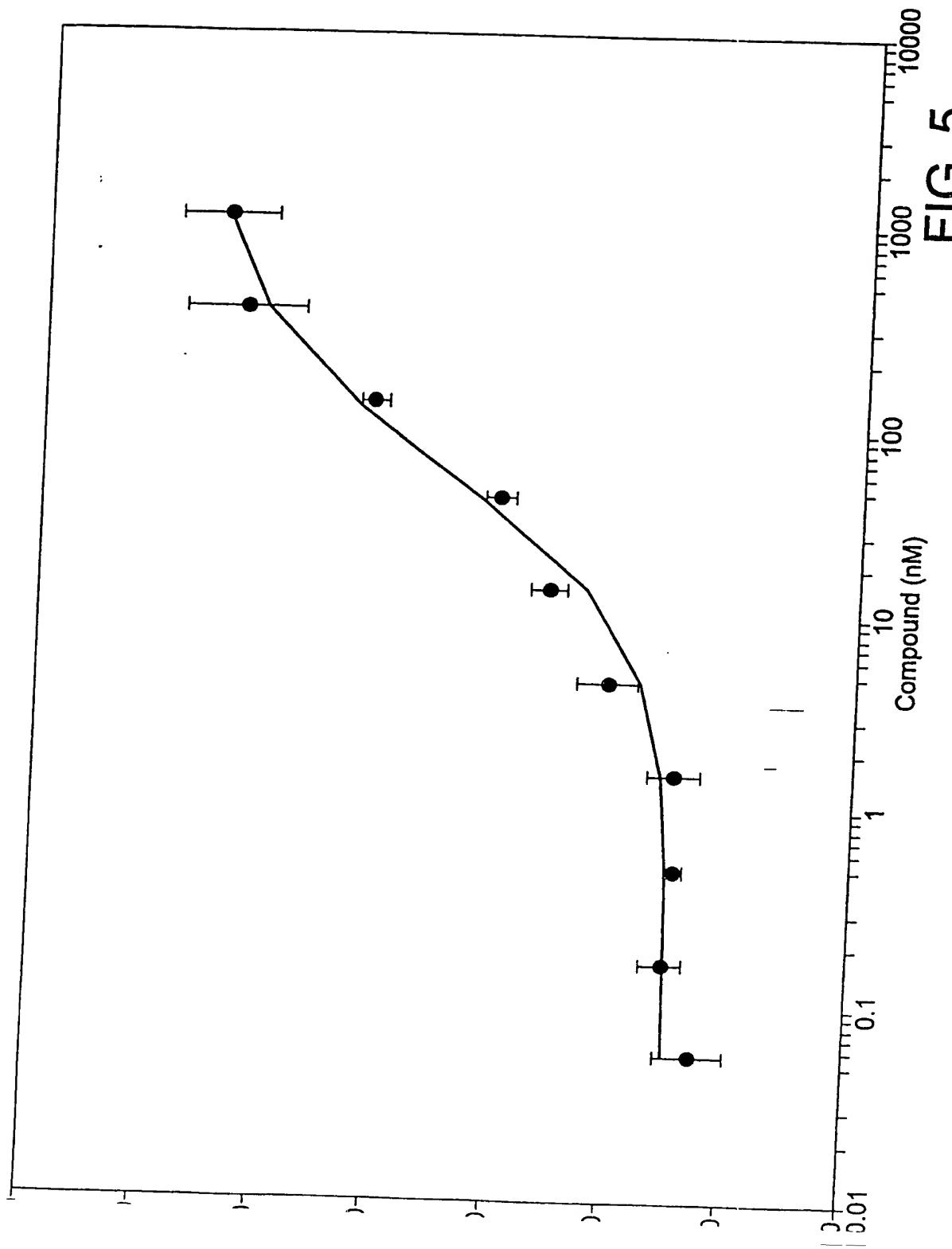
FIG. 3B.



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SEQUENCE LISTING

CPSSHSSLTERHKILHRL-QEGSPS (SEQ ID NO.: 1)
QEQLSPKKKENNALLRYLLDRDDPS (SEQ ID NO.: 2)
5 QEPVSPKKKENALLRYLLDKDDTKD (SEQ ID NO.:3)
GSTHGTSLKEKHKILHRLQDSSSPVD (SEQ ID NO.:4)
SNMHGSLLQEKHRLHKLLQNGNSPAE (SEQ ID NO.:5)
LXXLL (SEQ ID NO.:6)

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US99/30947

A. CLASSIFICATION OF SUBJECT MATTER

IPC(7) : A61K 31/42; C07K 14/00; G01N 33/53
US CL : 435/7.1, 7.8; 514/378; 530/324

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

U.S. : 435/7.1, 7.8; 514/378; 530/324

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)
WEST, MEDLINE, BIOSIS, EMBASE, CAPLUS, REGISTRY, GENBANK, SWISSPROT

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A Y	WO 96/21742 A1 (THE SALK INSTITUTE FOR BIOLOGICAL STUDIES) 18 July 1996 (18-7-96), pages 10-13.	8-11 ----- 17-33
P,Y	MAKISHIMA, M. et al. Identification of a Nuclear Receptor for Bile Acids. Science. 21 May 1999, Vol. 284, pages 1362-1365, especially page 1363, Figure 1.	1-7, 17-33
P,Y	PARKS, D.J. et al. Bile Acids: Natural Ligands for an Orphan Nuclear Receptor. Science. 21 May 1999, Vol. 284, pages 1365-1368, especially page 1366, Figure 2.	12-14 ----- -----

Further documents are listed in the continuation of Box C. See patent family annex.

Special categories of cited documents:	"T"	later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
A document defining the general state of the art which is not considered to be of particular relevance	"X"	document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
B earlier document published on or after the international filing date	"Y"	document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
L document which may throw doubt on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)	"&"	document member of the same patent family
O document referring to an oral disclosure, use, exhibition or other means		
P document published prior to the international filing date but later than the priority date claimed		

Date of the actual completion of the international search

31 MARCH 2000

Date of mailing of the international search report

02 MAY 2000

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US99/30947

C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	YAO, T-P. et al. The nuclear hormone receptor coactivator SRC-1 is a specific target of p300. Proc. Natl. Acad. Sci. USA. October 1996, Vol. 93, pages 10626-10631, especially page 10630, Figure 4.	1-4

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US99/30947

Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This international report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:

2. Claims Nos.:
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:

3. Claims Nos.: 34-35
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:

4. No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

The additional search fees were accompanied by the applicant's protest